GenCore version (c) 1993 - 2003

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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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    222222211111110
22222221176543210
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  of hits satisfying chosen parameters:
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Match
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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US-09-608-810A-2
US-08-975-080-11
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US-09-155-613A-46
US-09-171-482-7
US-08-931-222-42
US-08-933-813-422-42
US-08-933-833-422-42
US-08-934-223-42
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US-08-934-223-42
US-08-934-223-42
US-08-931-311-442C-1
US-08-931-311-311C-8
US-08-557-3098-46
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US-08-258-976-49
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2, Appli
11, Appl
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11, Appl
11, Appl
12, Appli
42, Appl
46, Appl
47, Appl
48, Appl
49, Appl
49, Appl
19, Appl
19, Appl
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US-08-975-080-11 US-08-975-080-11 ; Sequence 11, Application US/08975080 ; Patent No. 6245523 ; GENERAL INFORMATION: APPLICANT: Altieri, Dario C. TITLE OF INVENTION: SURVIVIN, A PROTEIN THAT INHIBITS TITLE OF INVENTION: CELLULAR APOPTOSIS, AND ITS MODULATION NUMBER OF SEQUENCES: 35 CORRESPONDENCE ADDRESS: ADDRESSEE: MORGAN, LEWIS & BOCKIUS LLP STREET: 1800 M Street, N.W. CITY: Washington STATE: D.C. COUNTRY: USA ZIP: 20036-5869 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk	Query Match 100.0%; Score 40; DB 4; Length 11; Best Local Similarity 100.0%; Pred. No. 0.034; Matches 8; Conservative 0; Mismatches 0; Indels 0; Qy 1 GSSFLSPE 8 Db 1 GSSFLSPE 8	RESULT 1 US-09-608-810A-2 US-09-608-810A-2 ; Sequence 2, Application US/09608810A ; Patent No. 6420521 ; GENERAL INFORMATION: APPLICANT: Sheppard, Paul O. ; APPLICANT: Jaspers, Stephen R. APPLICANT: Daisher, Theresa A. APPLICANT: Bishop, Paul D. ; TITLE OF INVENTION: SGIP PEPTIDES FILE REFERENCE: 99-51 ; CURRENT APPLICATION NUMBER: US/09/608,810A ; CURRENT APPLICATION NUMBER: 60/141,592 PRIOR APPLICATION NUMBER: 60/141,592 PRIOR FILLING DATE: 1999-06-30 ; PRIOR FILLING DATE: 1999-06-30 ; NUMBER OF SEQ ID NOS: 7 ; SOFTWARE: FastSEQ for Windows Version 3.0 ; SEQ ID NO 2 ; LENGTH: 11 ; ORGANISM: Homo sapiens	ALIGNMENTS	28 24 60.0 7 4 US-08-378-313-5 Sequence 29 24 60.0 12 1 US-08-196-989B-6 Sequence 30 24 60.0 12 2 US-08-196-989B-6 Sequence 31 24 60.0 13 2 US-08-637-759B-129 Sequence 32 24 60.0 13 3 US-08-637-759B-129 Sequence 32 24 60.0 13 4 US-09-201-945-129 Sequence 32 24 60.0 15 2 US-08-637-759B-129 Sequence 34 60.0 15 4 US-09-268-48B-3 Sequence 35 24 60.0 25 US-08-467-023-52 Sequence 36 24 60.0 20 3 US-08-467-023-52 Sequence 37 24 60.0 20 5 PCT-US95-06726-24 Sequence 39 24 60.0 21 1 US-08-471-780C-70 Sequence 40 24 60.0 21 1 US-08-471-780C-70 Sequence 41 24 60.0 21 2 US-08-467-10C-70 Sequence 42 60.0 21 2 US-08-467-10C-70 Sequence 43 24 60.0 21 US-08-468-7390-70 Sequence 43 24 60.0 29 4 US-09-311-311C-15 Sequence Sequence 50 29 4 US-09-311-311C-15 Sequence 50 29 20 25 20 25 20 25 20 25 20 25 25 25 25 25 25 25 25 25 25 25 25 25
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RESULT 5
US-09-171-482-7
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Fatent No. 6420120
GENERAL INFORMATION:
APPLICANT: Boulanger, Pierre
APPLICANT: Hong, Saw See
APPLICANT: Hong, Saw See
TITLE OF INVENTION: Uncie
TITLE OF INVENTION: Unce of a Polypeptide as Cell Receptor for Adenoviruses
FILE REFERENCE: 032751-036
CURRENT APPLICATION NUMBER: US/09/155,613A
CURRENT APPLICATION NUMBER: PCT/FR98/00184
PRIOR PILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: FR 97/01005
PRIOR PELING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
PRIOR FILING DATE: 1997-01-30
SPRIOR PELING DATE: 1997-01-30
SPRIOR PELING DATE: 1997-01-30
SOFTWARE: FastSEQ for Windows Version 4.0
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOOTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/975,080
FILING DATE: 20-MOV-1997
FILING DATE: 20-MOV-1997
FILING DATE: 20-MOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: AGIET, REALG G.
REGISTRATION NUMBER: 30,988
REFERENCE/DOCKET NUMBER: 30,988
REFERENCE/DOCKET NUMBER: 044574-5022-01-WO
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                           TELEFAX: 202-467-7176
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 45 amino acids
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ORGANISM: Artificial Sequence
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, OTHER INFORMATION: Phagotope
US-09-155-613A-46
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Best Local Similarity 71.4'
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; MOLECULE TYPE: protein US-08-975-080-11
                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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TOPOLOGY: li
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US-09-155-613A-46
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Sequence 7, Application US/09171482A

Sequence 7, Application US/09171482A

Patent No. 6184449

GENERAL INFORMATION:
A 1-AMINOSYCLOPROPANE-1-CARBOXYLATE SYNTHASE GENE FROM
TITLE OF INVENTION: ROSA TO CONTROL ETHYLENE LEVELS IN ROSES
FILE REPRENCE: TAGAWA-ROSE
CURRENT APPLICATION NUMBER: US/09/171,482A

CURRENT APPLICATION NUMBER: PCT/US97/17644, Published under WOS8/14465; US5,824,875

EARLIER FILING DATE: 1997-Sept-30, Published 1998-April-09; 1996-Oct-01

SOFTWARE: Word Perfect 6.1

FINGEN OF THE NOTE O
PATENT OF STATEMENT ON US/24194

BREERAL INFORMATION:

APPLICANT: RANU, RAJINDER S.

TITLE OF INVENTION: SYNTHAME GENES FROM PELARGONIUM TO CONTROL ETHYLENE LEVELS

TITLE OF INVENTION: SYNTHAME SINCH STANDIUMS

TITLE OF INVENTION: IN GERANIUMS

NUMBER OF SEQUENCES: 13

CONDESSEE: SANTAMGELO LAW OFFICES PC

STREET: 315 WEST OAK STREET, STE 701

CITY: FORT COLLINS

STATE: CO
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Pred. No. 53;
0; Mismatches 2; Indels
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Pred. No. 53;
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75.08;
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Best Local Similarity 75.0%;
Matches 6; Conservative (
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Best Local Similarity 75.0
Matches 6; Conservative
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MOLECULE TYPE: peptide
US-08-724-194-11
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; ORGANISM: Rosa kardinal
US-09-171-482-7
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STRANDEDNESS: si
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US-08-934-222-42

; Sequence 42, Application US/08934222

· Patent No. 5928895----
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Sequence 42, Application US/08933402
Patent No. 5948887
GENERAL INFORMATION:
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: U.S. FILING DATE: 23-APR-1993 ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/934,222 FILING DATE: 19-SEPT-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: EVANS, Herbert J.
APPLICANT: KINI, R. Manjunatha
TITLE OF INVENTION: Polypeptides That Include Conformation-
TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
TITLE OF INVENTION: Site
                                                                      APPLICANT: EVANS, Herbert J.
APPLICANT: KINI, R. Manjunatha
TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
TITLE OF INVENTION: Site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: U.S. FILING DATE: 29-OCT-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/5 FILING DATE: 03-MAY-1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                    CORRESPONDENCE ADDRESS:
                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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CITY: Washington
STATE: DC
                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 85.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Isacson, John P.
REGISTRATION NUMBER: 33,751
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ZIP: 20007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                               S SSTLSPE 11
                                                                                                                                                                                                                                                                                                                                   2 SSFLSPE 8
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Suite 500, 3000 K Street NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11 amino acids
l: Foley & Lardner
Suite 500, 3000 K Street NW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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Pred. No. 24;
                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                        Length 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 42,
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INFORMATION FOR SEQ ID NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 42, Application US/09207621
Patent No. 5952465
GENERAL INFORMATION:
APPLICANT: EVANS, Herbert J.
APPLICANT: KINI, R. Manjunatha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/953/,
FILING DATE: 19-SEPT-1997
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/532,818
                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Polypeptides That Include Conformation-
TITLE OF INVENTION: Constraining Groups Which Flank A Proten-Protein Interaction
NUMBER OF SEQUENCES: 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: U.S. FILING DATE: 23-APR-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 29-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 08/532,010
APPLICATION NUMBER: 08/532,010
FILING DATE: 03-MAY-1996
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: U.S. 08/143,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                 FILING DATE: 21-APR-1994 PRIOR APPLICATION DATA:
                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                          STREET: Suite 500, 3000 K Street NW CITY: Washington STATE: DC COUNTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Isacson, John P. REGISTRATION NUMBER: 33,751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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ZIP: 20007
                                                     APPLICATION NUMBER:
                                                                  APPLICATION NUMBER: 08/5 FILING DATE: 03-MAY-1996
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                                                                                                                            FILING DATE:
                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                             ZIP: 20007
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es 6; Conserv
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)GY: linear
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85.7%;
                                                   PCT/US94/04294
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U.S. 08/143,364
                                                                                     08/532,818
                                                                                                                                           US/09/207,621
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Pred. No. 24;
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Sequence 42, Application US/08934224

| Patent No. 6100044
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: EVANS, Herbert J.
| APPLICANT: KINI, R. Manjunatha
| TITLE OF INVENTION: Polypeptides That Include Conformation-
| TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
| TITLE OF INVENTION: Site
                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: EVANS, Herbert J.
TITLE OF INVENTION: Polypeptides That Include Conformation-
TITLE OF INVENTION: Constraining Groups Which Flank A Proten-Protein Interaction
NUMBER OF SEQUENCES: 153
ADDRESSEE: Folly & Lardner
STREET: Suite 500, 3000 K Street NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPEATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/231,797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 26; DB 3;
Pred. No. 24;
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APPLICATION NUMBER: 08/532,818
FILING DATE: 03-MAY-1996
APPLICATION NUMBER: PCT/US94/04294
FILING DATE: 21-APR-1994
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
FILING DATE: 29-0CT-1993
PRIOR APPLICATION NUMBER: 0.S. 08/143,364
FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: 18acson, John P.
REGISTRATION NUMBER: 040433/0148
INFORMATION FOR SEQ ID NO: 42:
SECRENCE/DOCKET NUMBER: 040433/0148
INFORMATION FOR SEQ ID NO: 42:
SECRENCE/DOCKET NUMBER: 040433/0148
                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Foley & Lardner
STREET: Suite 500, 3000 K Street NW
CITY: Washington
                                                                                                                                                                           US-09-231-797-42; Sequence 42, Application US/09231797; Patent No. 6084066
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85.7%;
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GY: linear
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Best Local Similarity
                                                                                   SSTLSPE 11
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                                    SSFLSPE 8
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US-08-934-224-42
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Sequence 42, Application US/08532818
Sequence 42, Application US/08532818
Setent No. 5965698
GENERAL INFORMATION:
APPLICANT: EVANS. Herbert J.
APPLICANT: KINI, R. Manjunatha
TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
TITLE OF INVENTION: Size
CORRESPONDENCE ADDRESS:
ADDRESSBE: Foley & Lardner
STREET: Suite 500, 3000 K Street NW
CITY: Washington
STRATE: DC
COURTRY: USA
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Pred. No. 24;
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85.7%; Pred. No. 24;
tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/532,818
FILING DATE: 03-MAY-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US. 08/143,364
FILING DATE: 21-APR-1994
PRIOR APPLICATION NUMBER: US. 08/143,364
FILING DATE: 29-OCT-1993
PRIOR APPLICATION NUMBER: US. 08/143,364
FILING DATE: 23-APR-1993
ATPONEY/AGENT INFORMATION:
ANAME: 10.50.007
                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
FILING DATE: 29-OCT-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. 08/051,741
FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: ISAGSON, John P.
REGISTRATION NUMBER: 33,751
REFERRINGE/DOCKET NUMBER: 040433/0148
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: amino acid
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REGISTRATION NUMBER: 33,751
REFERNCE/DOCKET NUMBER: 040433/0148
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
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Matches 6; Conservative
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3Y: linear
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Best Local Similarity
Matches 6; Conserv
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US-08-933-843-42
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/934,224
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/934,224
FILING DATE: 03-WAY-1996
APPLICATION NUMBER: 00/532,818
FILING DATE: 03-WAY-1996
APPLICATION NUMBER: PCT/US94/04294
PRIOR APPLICATION DATA:
APPLICATION DATA: 06/14204
PRIOR APPLICATION DATA: 06/14204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: EVANS, Herbert J.
APPLICANT: KINI, R. Manjunatha
TITLE OF INVENTION: Polypeptides That Include Conformation-
TITLE OF INVENTION: Constraining Groups Which Flank A Protein-Protein Interaction
TITLE OF INVENTION: Site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: IBACSON, John P.
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 4
               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/933,843
FILING DATE: 19-SEPT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: U.S. 08/143,364
FILING DATE: 29-OCT-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. 08/051,741
FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
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                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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RIOR APPLICATION DATA:
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CITY: Washington
STATE: DC
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TOPOLOGY: linear
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                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                CITY: Washington
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Suite 500, 3000 K Street NW
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Suite 500, 3000 K Street NW
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85.7%;
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RESULT 13
US-08-934-223-42
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Patent No. 61
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REGISTRATION NUMBER: 33,75
REFERENCE/DOCKET NUMBER: 0.
INFORMATION FOR SEQ ID NO: 42:
                                   NAME: ISACEON, JOhn P.
REGISTRATION NUMBER: 33,75
REFERENCE/DOCKET NUMBER: 0
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/934,22
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MEDIUM TYPE: Floppy of COMPUTER: IBM PC com
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LENGTH: 11 amino acids
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                                                                                                      FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US94/04294
FILING DATE: 21-APR-1994
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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TITLE OF INVENTION:
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Local Similarity 85.7%;
les 6; Conservative
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OPERATING SYSTEM: PC-DOS/MS-DOS
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FILING DATE: 29-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
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                                                                                                                                                                      APPLICATION NUMBER: U.S. 08/143,364 FILING DATE: 29-OCT-1993
                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/01
FILING DATE: 03-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20007
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Suite 500, 3000 K Street NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
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Constraining Groups Which Flank A Protein-Protein Interaction
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0; Mismatches
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                                                   040433/0148
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APPLICANT: K. KIMURA, et al.
TITLE OF INVENTION: NOVEL PROLYL ENDOPEPTIDASE
TITLE OF INVENTION: INHIBITORS SNA-115 AND SNA-115TILE OF INVENTION: PRODUCTIVE STRAIN THEREOF
INTILE OF INVENTION: PRODUCTIVE STRAIN THEREOF
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: BUTGESS:
ADDRESSEE: 370 Lexington Avenue, Suite 2105
CITY: NEW YORK
STREET: NEW YORK
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COUNTRY: UNITED STATES OF AMERICA
ZIP: 10017
COMPUTER READABLE FORM:
MEDIUM TYPE: 5 1/4 inch diskette
COMPUTER: Ev. LIMITED SYSTEM 200
OPERATING SYSTEM: DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/977,444C
FILING DATE: 17 - FEBRUARY 1993
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 17,906
REGISTRATION NUMBER: 17,906
REFERENCE/DOCKET NUMBER: 17,906
REFEREN
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Job time : 12.3636 secs
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TOPOLOGY: CYCLIC OF LINEAR
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity
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Sequence 42, Application US/09413492

Sequence 42, Application US/09413492

Sequence 42, Application US/09413492

Sequence 42, Application US/09413492

SENERAL INFORMATION:

APPLICANT: EVANS, Herbert J.

APPLICANT: EVANS, Herbert J.

APPLICANT: EVANS, Herbert J.

TITLE OF INVENTION: Oolypeptides That Include Conformation-
TITLE OF INVENTION: Site

NUMBER OF SEQUENCES: 153

CORRESPONDENCE ADDRESS:

ADDRESSEE: Foley & Lardner

STREET: Suite 500, 3000 K Street NW

CONTY: Mashington
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                                                                                                                   65.0%; Score 26; DB 4; Length 11; 85.7%; Pred. No. 24; 1; Indels ive 0; Mismatches 1; Indels
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Pred. No. 24;
0; Mismatches 1; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/413,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/532,818
APPLICATION NUMBER: 08/532,818
APPLICATION NUMBER: PCT/US94/04294
FILING DATE: 03-MAY-1996
FILING DATE: 21-APR-1994
PRIOR APPLICATION DATA:
APPLICATION DATA: 0.S. 08/143,364
FILING DATE: 22-OCT-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. 08/051,741
FILING DATE: 23-APR-199
ATTONNEY/AGENT INFORMATION:
NAME: Isaccon, John P.
REGISTRATION NUMBER: 33,751
REFERENCE/DOCKET NUMBER: 33,751
REFERENCE/DOCKET NUMBER: 040433/0148
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHRACTERISTICS:
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Best Local Similarity 85.7%;
Matches 6; Conservative (
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Matches 6; Conservative
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TYPE: amino acid
TOPOLOGY: linear
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                 ; TOPOLOGY:
US-08-934-223-42
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STATE: DC
COUNTRY:
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US-07-977-444C-1
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length:
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1. /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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          GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd
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US-09-201-945-29
US-09-901-12B-1
PCT-US95-06726-24
US-07-942-245-383
US-08-724-194-11
US-09-171-482-7
5198359-8
5284931-17
544976-11
US-08-618-696-13
US-09-033-753-13
5438119-11
US-08-8189-841B-54
US-08-8189-841B-54
US-08-818-841B-644
US-08-444-818-644
US-08-444-818-645
US-08-444-818-646
US-09-033-731-7
US-08-350-260A-435
US-08-925-002-52
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Sequence 2, Appli
Sequence 129, App
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Sequence 124, Appli
Sequence 24, Appli
Sequence 17, Appli
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Sequence 13, Appli
Patent No. 5284931
Patent No. 5284931
Patent No. 54849756
Sequence 13, Appli
Sequence 54, Appli
Sequence 54, Appli
Sequence 64, Appli
Sequence 642, Appli
Sequence 643, Appli
Sequence 644, Appli
Sequence 645, Appli
Sequence 645, Appli
Sequence 646, Appli
Sequence 646, Appli
Sequence 647, Appli
Sequence 647, Appli
Sequence 648, Appli
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Sequence 648, Appli
Sequence 52, Appli
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RESULT 2 US-08-637-759 ; Sequence 12 ; Patent No. ; GENERAL IN, ; APPLICAN APPLICAN OCORRESPO ADDRES STREET STREET CITY: ; STATE: ; COUNTR ZIP: COMPUTER MEDIUM	Query Matc Best Local Matches Qy 1 GS Db 1 GS	RESULT 1 US-09-608-810 Sequence 2, Patent No. GENERAL INF APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT INTE OF I FILE REFER CURRENT AP CURRENT AP CURRENT APPL PRIOR APPL PRIOR APPL PRIOR APPL PRIOR APPL PRIOR FILI NUMBER OF SOFTWARE: SEQ ID NO 2 LENGTH: 1 TYPE: PRI ORGANISM: US-09-608-810	222CCCCCCCCCCCA444A800000000000000000000000
OF TRY	Match Jocal Simi ss 4; 1 GSSF 4 1 GSSF 4	8-810A-2 NO. 6420521 L INFORMATION CANT: Sheppa CANT: Jasper CANT: Deishe CANT: Bishop OF INVENTION REFERENCE: 9 NT APPLICATION PILLING DATE APPLICATION FILING DATE APPLICATION FILING DATE APPLICATION FILING DATE APPLICATION TO SEQ ID N ARE: FastSEQ NO 2 TH: 11 : PRT NISM: Homo sa 8-810A-2	111111111111111111111111111111111111111
HUNGHROHROKUM	100.0 nilarity 100.0 Conservative	B-810A-2 INC 2, Application US/0 INC 2, Application US/0 INC 6420521 IL INCORMATION: ICANT: Sheppard, Paul ICANT: Sheppard, Stephe ICANT: Jaspers, Stephe ICANT: Bishop, Paul D: SOF INVENTION: SGIP P REFERENCE: 99-51 SOF INVENTION NUMBER: ANT APPLICATION NUMBER: APPLICATION NUMBER: R FILING DATE: 2000- R APPLICATION NUMBER: R FILING DATE: 199-06 R OF SEQ ID NOS: 7 AARE: FastSEQ for Wind NO 2 STH: 11 SIPRT ANISM: Homo sapiens 18-810A-2	88888888888888888888888888888888888888
icon icon icon icon icon icon icon icon	100. 100. rative	ication US/09608810A [21] [ION: [Dard, Paul O.] [Ders, Stephen R.] [Sher, Theresa A.] [Hop, Paul D.] [ION: SGIP PEPTIDES [199-51] [ITION HUMBER: US/09/6 [DATE: 2000-06-30 [ON NUMBER: 60/141,5] [TE: 199-06-30 [D NOS: 79.66-30] [D	110 110 110 111 111 113 113 113
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100.0%; Score 20; DB 3; Length 13; Conservative 0; Mismatches 0; Indels
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100.0%; Score 20; DB 4; Length 13;
Best Local Similarity 100.0%; Pred. No. 56;
Matches 4; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Alauta
STATE: Georgia
COUNTRY: USA
ZIP: 30309-3460
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BM PC Compatible
OMPRATING SYSTEM: PC-DOS/MS-DOS
CURRENT PAPLICATION DATA:
APPLICATION NUMBER: US/09/201,945
FILING DATE: CLASSIFICATION:
PREDICATION NUMBER: 08/637,759
FILING DATE: CLASSIFICATION:
APPLICATION NUMBER: 08/637,759
FILING DATE: CLASSIFICATION:
REFERENCE/DOCKET NUMBER: 31,284
REFERENCE/DOCKET NUMBER: 31,284
REFERENCE/DOCKET NUMBER: 31,284
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-201-945-129
Sequence 129, Application US/09201945
Fetent No. 6342215
GENERAL INFORMATION:
APPLICANT: David William Holden
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 501
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
                                                                                                                  LENGTH: 13 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLLGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS: LENGTH: 13 amino acids
                                                  INFORMATION FOR SEQ ID NO. SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
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                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 4; Conserv
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STRANDEDNESS: si
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100.0%; Score 20; DB 2; Length 13;
Best Local Similarity 100.0%; Pred. No. 56;
Matches 4; Conservative 0; Mismatches 0; Indels
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/637,759B
FILING DATE: 03-MAY-1996
CLASSIFCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB95/02875
FILING DATE: 1-DEC-1995
ATTONEY/AGENT INFORMATION:
NAME: Paber, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: 31,284
REGISTRATION OFFER STO ID NO: 129:
SEQUENCE CHARACTERISTICS:
LEMGTH: 13 amino acids
TYPE: amino acid
STRANDEDNESS: single
TYPE: Amino acid
STRANDEDNESS: single
MUDECTILE TWOST.
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1 US-08-871-355A-129

2 Sequence 129, Application US/08871355A

Patent N. 6015669

3 GENERAL INFORMATION:
Patent Inchritication of Genes
CORRESPONDENCE ADDRESS:
ADDRESSE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
CITY: Antlanta
STATE: Georgia
COMPUTER: Georgia
COMPUTER: Eloppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MM-DOS
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MM-DOS
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MM-DOS
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MM-DOS
SOURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/GB95/02875
FILING DATE: 11-DEC-1995
CLASSIFICATION: 435
FILING DATE: 11-DEC-1995
CLASSIFICATION: 435
FILING DATE: 11-DEC-1995
CLASSIFICATION NUMBER: RPMS 101 CON
TELECOMMUNICATION NUMBER: RPMS 101 CON

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (301) 977-0847
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDENNESS: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 74;
Matches 4; Conservative 0; Mismatches
                                                                                                                                                                                                                                Sequence 24, Application PC/TUS9506726 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                               TITLE OF INVENTION: Ligands for Induction of Antigen Specific NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
ADDRESSE: LAHIVE & COCKFIELD
STREET: 60 STATE STREET, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: not releval MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/478,591
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Gormley, Mary E.
REGISTRATION NUMBER: 34,409
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 258-5200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:
                                                           STREET: 60 ST
CITY: BOSTON
STATE: MASSAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT: Hanna, Micheal G.
PPLICANT: Haspel, Martin V.
PPLICANT: Hoover, Herbert C.
ITLE OF INVENTION: Tumor Associated Epitopes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Akzo No. 5951985el Patent Department STREET: 1300 Piccard Drive, Suite 206 CITY: Rockville STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/960,128 FILING DATE: CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
ZIP: 20850
                      COUNTRY: USA
ZIP: 02109-1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1, Application US/08960128
5, 5951985
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Y: USA
                                                           MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Butler, Sandra M.
Pomato, Nicholas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   not relevant
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TYPE: amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCT-US95-06726-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Sest Local Similarity 100.0%; Matches 4; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: HP 9000/700 Workstation

OPERATING SYSTEM: UNIX

SOFTWARE: In house

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/942,245
TELEX: 6491103
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
APPLICANT:
                                                                          FILING DATE: 09-SEP-1992
CLASSIFICATION: 530
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: RPI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Sughrue, Mion, Zinn, Macpeak & Seas
STREET: 2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06726
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: GUILD, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                             TELEPHONE: (202) 293-70
TELEFAX: (202) 293-7860
                                                                                                                                                                                                                                                                                            CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC
SOFTWARE: ASCII text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 03 JUICLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
COMPUTER: IB
                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GSSF 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                383, Application US/07942245
                                                                                                                                                                                                                                                          20037-3202
                                                                                                                                                                                                                                                                                        D.C.
                                                                                                                                                                                                                                                                       United States
                                                                                                                                                                                                                                                                                                                                                                                                                                           ROGUSKA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEARLE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEDERSEN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide
internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03 JUNE 1994
                                                                                                                                                                                                                                                                                                                                                                                          Stephen M.J.
Anthony R.
Michael A.
Braydon C.
SURFACE RESIDUE VENEERING OF RODENT
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jan T
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Gaps

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; CURRENT FILING DATE: 1998-10-19
; EARLIER APPLICATION NUMBER: PCT/US97/17644, Published under WO98/14465; US5,824,875
; EARLIER FILING DATE: 1997-Sept-30, Published 1998-April-09; 1996-Oct-01
; SOGTWARE: Word Perfect 6.1
; SOGTWARE: Word Perfect 6.1
; ENGTH 35
; TYPE: PRT
; ORGANISM: Rosa kardinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5449756-11
;Patent No. 5449756
; APPLICANT: TANIGUCHI, TADATSUGU;HATAKEYAMA, MASANORI;MINAMOTO,
;SEJIRO;KONO, TAKESHI;DOI, TAKESHI;MIYASAKA, MASAYUKI;TSUDO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 11
5284931-17
FARENT NO 5284931
FARENT NO 5284931
TITLE OF INVENTION;
FILE OF INVENTION:
FILE OF INVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: TANIGUCHI, TADATSUGU;JATAKEYAMA, MASANORI;
MINAMOTO, SEJIRO;KONO, TAKESHI;DOI, TAKESHI;MIYASAKA, MASAYUKI;
;TSUDO, MITSURU;KARASUYWA, HAJIME
TITLE OF INVENTION: RECOMBINANT PROTEIN RECEPTOR FOR IL-2
NUMBER OF SEQUENCES: 9
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/487,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 20; DB 4; Length 35; 100.0%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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75.0%; Pred. No. 1.9e+05;
Live 1; Mismatches 0; Indels
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Best Local Similarity 75.0%; Pred. No. 1.9e+05;
Matches 3; Conservative 1; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 4; Conservative
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; LENGTH: 4
5198359-8
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1 GASF 4
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;Patent No.
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| Sequence 11, Application US/08724194 |
| Sequence 11, Application US/08724194 |
| Sequence 11, Application US/08724194 |
| Patent No. 5824875 |
| GENERAL INFORMATION: TO SEQUENCE SEQUENCE SEQUENCE SEQUENCE SEQUENCE SEQUENCE SEQUENCE SEQUENCES SWITHASE GENES FROM PELARGONIUM TO CONTROL ETHYLENE LEVELS NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESSE: SANTANGELO LAW OFFICES PC STREET: 315 WEST OAK STREET, STE 701 |
| STREET: CO COUNTRY: USA STREET STE NOW NOT COLLINS STATE: CO COUNTRY: USA STREET STE NOT COLLINS STATE: CO COUNTRY: USA CONTRY: USA CONTRY: USA CONTRY: USA COUNTRY: USA CO
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Patent No. 618449
GENERAL INFORMATION:
APPLICANT: REMINER SIGNATION:
TITLE OF INVENTION: A 1-AMINOCYCLOPROPANE-1-CARBOXYLATE SYNTHASE GENE FROM TITLE OF INVENTION: ROSA TO CONTROL ETHYLENE LEVELS IN ROSES
FILE REFERENCE: TAGAMA-ROSE
CURRENT APPLICATION NUMBER: US/09/171,482A
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                                                                                                                                                                                                                                          Length 26;
                                                                                                                                                                                     Score 20; DB 1; Lengtn 20
Pred. No. 1.1e+02;
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100.0%; Pred. No. 1.58+02;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PATCHTION DATA:
SOFTWARE: PATCHTION DATA:
APPLICATION NUMBER: US/08/724,194
FILING DATE: 01-0CT-1996
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: SANTANGELO, LUKE
REGISTRATION NUMBER: 31,997
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 amino acids
                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                         Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 4; Conservative 0
, TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-07-942-245-383
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Best Local Similarity 100.
Matches 4; Conservative
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TOPOLOGY: linear
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US-09-171-482-7
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RESULT 13
US-08-618-696-13
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FILING DATE: 08-APR-1992
APPLICATION NUMBER: 487,059
FILING DATE: 05-MAR-1990
SEQ ID NO:11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MITSURU; KARASUYAMA, HAJIME
; TITLE OF INVENTION: RECOMBINANT PROTEIN RECEPTOR FOR IL-2
; NUMBER OF SEQUENCES: 12
; CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 13, Application US/08618696
Patent No. 5861475
GENERAL INFORMATION:
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Best Local Similarity
Matches 3; Conser
                                                            Query Match
                               Matches
                                                                                                                                                                                                                                                                                         MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDDERFECT 5.1
CURRENT APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION DATA:
APPLICATION UMBER: 07/995,269
FILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
NAMS: PARKER, DAVID L.
REGISTRATION NUMBER: 3,165
REFERENCE/DOCTET NUMBER: 3,165
                                                                                                                                                                          TELEX: NOT APPLICABLE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/88,592 FILING DATE: 9-JUL-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: UO TELECOMMUNICATION INFORMATION: TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: ARNO
STREET: P.O. BO
CITY: HOUSTON
STATE: TEXAS
                             Local Similarity nes 3; Conserv
                                                                                                              TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
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1 GSSF 4
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P.O. BOX 4433
                                                                                                                                                              5 amino acid residues
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                                                                                                          linear
                               Conservative
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                                              85.0%;
75.0%;
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75.0%;
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3ER: UOAB:002/PAR
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                  Score 17; DB 2; Length 5,
Pred. No. 1.9e+05;
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Pred. No. 1.9e+05;
1; Mismatches 0
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RESULT 15
5438119-11
;Patent No.
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US-09-033-753-13
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Best Local Similarity
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NUMBER OF SEQUENCES: 16
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/07/98
FILING DATE: 25-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 525,899
FILING DATE: 18-MAY-1990
APPLICATION NUMBER: 189,318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: NOT APPLICABLE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/618
FILING DATE: 20-MAR-1996
APPLICATION NUMBER: 07/995
PILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09
                                                                                                                                                         APPLICANT: Rutter, William; Santi, Daniel TITLE OF INVENTION: METHOD OF OBTAINING A PEPTIDE WITH DESIRED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 5 amino acid residues
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                     Conservative
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                      85.0%;
75.0%;
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                                                                                                                                                                                                                                                                                                                                                   Score 17; DB 3;
Pred. No. 1.9e+05,
1; Mismatches (
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; FILING DATE: 02-MAY-1988
; SEQ ID NO:11:
; LENGTH: 5
5438119-11

Query Match
Best Local Similarity 75.0%; Pred. No. 1.9e+05;
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps
Qy 1 GSSF 4

Db 1 GSAF 4
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Search completed: January 10, 2003, 15:59:54 Job time : 8.18182 secs

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Minimum DB
Maximum DB
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No.
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
         Pred. No. is th score greater than and is derived
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eq
       length: 0 length: 50
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                       is the number of results predicted by chance to have a ster than or equal to the score of the result being printed, rived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/FCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
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Gapop 10.0 ,
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    US-08-875-277A-13
US-08-836-922-18
5182210-22
5182210-16
US-08-159-339A-824
US-08-159-339A-824
US-09-385-7408-17
US-09-385-7408-19
US-09-385-7408-19
US-09-385-7408-19
US-09-385-7408-19
US-08-548-974-24
US-08-548-974-25
US-08-548-974-17
US-08-93-712-2
US-08-93-712-2
US-08-94-957-107
US-08-94-957-107
US-08-974-775-46
US-08-974-755-644-1
US-08-950-953-644-1
US-08-950-953-644-1
US-08-950-953-3321
US-08-950-923-3321
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19.038 Million cell updates/sec
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Sequence 13, Appl
Sequence 18, Appl
Patent No. 5182210
Patent No. 5182210
Sequence 824, App
Sequence 854, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 15, Appl
Sequence 24, Appl
Sequence 27, Appl
Sequence 17, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 17, Appl
Sequence 18, Appl
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RESULT 1 US-08-875-277A-: Sequence 13,: GENERAL INFOLANT: APPLICANT: APPLICANT: INTIES OF 11 INUMBER OF: CORRESPONDI APPLICAT: COMPUTER R. SOFTWARE CURRENT APPLICAT: FILING D. CLASSIFIL PRIOR APPLICAT: FILING D. A	00000000000000000000000000000000000000
277A-13 277A-13 16,79 16,70 10	226 226 226 226 227 227 227 227 227 227
S/088 DAVID TOPHE R J. TANT TANT TANT TOPHE R J. TOPHE	39 5 PCT 110 2 US- 113 3 US- 113 3 US- 114 US- 20 2 US- 21 4 US- 22 4 US- 23 4 US- 24 US- 25 US- 26 US- 27 US- 28 US- 28 US- 28 US- 29 US- 20 US- 20 US- 20 US- 20 US- 21 US- 21 US- 22 US- 23 US- 24 US- 25 US- 26 US- 27 US- 27 US- 28 US- 28 US- 28 US- 29 US- 29 US- 20
ALIGNMENTS 875277A D J. ER R. LUCIFERASES LUCIFERASES PMS-DOS FMS-DOS FMS-	08-535-31 -US94-05 08-536-5 08-537-1 08-637-1 08-871-3 09-201-9 09-143-1 08-811-4 08-811-4 08-813-3 09-220-5 09-220-5 09-220-5 08-486-0 08-486-0
n #1.30	51-1 468-1 97-171 59-129 55A-129 45-129 44-145 92-145 92-145 92-115 92-119 93-13 98-12 98-12 98-12 98-13 98-12 98-12 98-12 98-13 98-14
	Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 2 Sequence 2 Sequence 2 Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 2 Sequence 3 Sequence 3 Sequence 3 Sequence 8 Sequence 8 Sequence 8 Sequence 9 Sequence 7
•	1, Appli 1, Appli 1, Appli 129, App 129, App 129, App 129, App 129, App 145, Appli 145, Appli 19, Appli 19, Appli 19, Appli 19, Appli 11, Appli 11, Appli 11, Appli 12, Appli 13, Appli 13, Appli 14, Appli 18, Appli 19, Appli 19

Query Match Best Local Similarity

58.0%;

Score 29; DB 4; Pred. No. 13;

Length

Matches

. 4

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Gaps

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3; Indels

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Score 28; DB 6; Length 32;
Pred. No. 33;
1; Mismatches 3; Indels
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US-08-159-339A-824

Sequence 824, Application US/08159339A
Fatent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Gete, Alessandro
ITILE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two send and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                    Patent No. 5182210

APPLICANT: BINNS, MATTHEW M.; BOURSNELL, MICHAEL E.G.;
CAMPBELL, JOAN I.A.; TOMLES, FIONA M.
TITLE OF INVENTION: FOWLFOX VIRUS PROMOTERS
NUMBER OF SEQUENCES: 22
CURRENT APPLICATION DATA:
APPLICATION DATA:

RELING DATE: 21-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: DEM COMPATIBLE
OPERATING SYSTEM: DOS
SOFTWARE: FASESED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/027,746
FILING DATE: 06-AUG-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 106-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.0%; Score 28; DB 6; 60.0%; Pred. No. 53; tive 1; Mismatches
                                                                                                          56.0%;
                                                                                                 Query Match
Best Local Similarity 60.0
Matches 6; Conservative
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Best Local Similarity 60.0
                                                                                                                                                                                       1 GSSFAKLOPR 10
                                                                                                                                                                                                                              5 GSSRVDLOPK 14
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                                        ; LENGTH: 32
5182210-22
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; LENGTH: 50
5182210-16
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                  SEQ ID NO:22:
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;Patent No.
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WELLS
WENDLEART: INNES PROUDFOOT, AMANDA ELIZABETH
MAPPLICANT: WELLS, TIMOTHY NIGEL CAMANDA ELIZABETH
MAPPLICANT: WELLS, TIMOTHY NIGEL CAMANDA ELIZABETH
TITLE OF INVENTION: COMPOSITIONS COMPRISING IT FOR TREATMENT OF INFLAMMATION
NUMBER OF SEQUENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CONTRY: ALLINGTON
STATE: VIRGINA
CONTRY: ALLINGTON

STATE: VIRGINA
CONTRY: LES PREATE FORM:
MEDIUM TYPE: FORM:
MEDIUM TYPE: FLODE/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: GB 9424835.8
FILING DATE: 16-UNN-1995
ATTORNEY/ARREIT INFORMATION:
APPLICATION NUMBER: GB 9512319.6
FILING DATE: 16-UNN-1995
ATTORNEY/ARREIT INFORMATION:
APPLICATION WINDER: GB 9512319.6
FILING DATE: 16-UNN-1995
ATTORNEY/ARREIT INFORMATION:
APPLICATION WINDER: MADAY TIMENDER:
APPLICATION WINDER: GB 94248315.9
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Pred. No. 17;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FALENCE DE STATEM M.; BOURSNELL, MICHAEL E.G.;
CAMPBELL, JOAN I.A.; TOMLEY, FIONA M.; TITLE OF INVENTION: FOWLPOX VIRUS PROMOTERS
NUMBER OF SEQUENCES: 22
CURRENT APPLICATION DATA;
APPLICATION NUMBER: US/08/469,608
FILING DATE: 21-OCT-1988
      1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: WILSON, MARY J.

REGISTRATION UNDHER: 32,955
REFERENCE/DOCKET NUMBER: 1430
TELECOMMUNICATION INFORMATION:
TELEFAX: (703) 816-401
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               56.0%;
6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 60.0
Matches 6; Conservative
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                                      1 GSSFAKLOPR 10
                                                                        8 GSSTVDLQPK 17
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                                                                                                                                              RESULT 2
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Length 50; 3; Indels

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RESULT 6
US-08-159-339A-854
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                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FeatSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION 424
PRIOR APPLICATION DATA:
TENNIC DATE: 115 07/926,666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                       APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/ACENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
MEDIUM TYPE: IBM Compatible
Compatible
DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessand
APPLICANT: Celis, Esteban
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 9 amino acids
                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: Two Embarca
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415) 576-020
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 32,762
                           STRANDEDNESS:
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10. 60371
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Celis, Esteban
                                                                                                                                           (415) 576-0300
linear
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Pred. No. 1.9e+05;
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FILE REFERENCE: NYU-011
CURRENT APPLICATION NUMBER: US/09/001,984C
CURRENT FILING DATE: 1997-12-31
PRIOR APPLICATION NUMBER: 60/034,003
PRIOR FILING DATE: 1996-12-31
NUMBER OF SEQ ID NOS: 106
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 72
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity bo...
Thehes 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vequence 17, Application US/09385740B
Patent No. 6348320
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/335,098
PRIOR FILING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: US 60/089,823
PRIOR FILING DATE: 1998-06-19
NUMBER OF SEQ ID NOS: 31
ORGANISM: Mycobacterium tuberculosis strain H37Rv-09-001-984C-72
                                                                                                                                                                                                                    APPLICANT: Laal, Suman
APPLICANT: Zolla-Pazner, Susan
APPLICANT: Belisle, John T
TITLE OF INVENTION: EARLY DETECTION OF MYCOBACTERIAL DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/385,740B CURRENT FILING DATE: 1999-08-30 PRIOR APPLICATION NUMBER: US 60/142,274 PRIOR FILING DATE: 1999-07-02 PRIOR APPLICATION NUMBER: US 60/141,574 PRIOR FILING DATE: 1999-06-29 PRIOR FILING DATE: 1999-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Eyre, David
TITLE OF INVENTION: CARTILAGE RESORPTION ASSAYS
FILE REFERENCE: WROS-1-14269
                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (1)..(18)
OTHER INFORMATION: syn- corr. to C-terminal telopeptide seq.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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66.7%;
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    Mismatches

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Pred. No.
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Pred. No.
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29;
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NAME/KEY: PEPTIDE

LOCATION: (1)...(21)

OTHER INFORMATION: syn- corr. to C-terminal telopeptide seq. of hu. type II collage.

US-09-385-740B-19
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Sequence 24, Application US/08548974

Sequence 24, Application US/08548974

Setent No. 5939529

GENERAL INFORMATION:

APPLICANT: Potenmpa, Lawrence A

TITLE OF INVENTION: Production Of Megakaryocytes And Thrombocytes

INDER OF SEQUENCES: 24

CORRESPONDENCE ADDRESS:

ADDRESSEE: BRINKS HOFER GILSON & LIONE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      54.0%; Score 27; DB 4; Length 21; 66.7%; Pred. No. 34;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPBERTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/548,974
FILING DATE: 27-OCT-1995
CLASSIFICATION: 435
                                                                                                                       GENERAL INCORMATION:
APPLICANT: Eyre, David
TILLE OF INVENTION: CARTLIAGE RESORPTION ASSAYS
FILE REPRENCE: WROS-1-1469
CURRENT APPLICATION NUMBER: US/09/385,740B
CURRENT FILING DATE: 1999-08-30
PRIOR PELICATION NUMBER: US 60/142,274
PRIOR PELICATION NUMBER: US 60/142,274
PRIOR PELICATION NUMBER: US 60/141,574
PRIOR PELING DATE: 1999-06-29
PRIOR FILING DATE: 1999-06-19
PRIOR FILING DATE: 1999-06-17
PRIOR PELING DATE: 1998-06-19
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTING DATE: 1998-06-19
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTING DATE: 1998-06-19
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CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/202,033
FILING DATE: 23-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Martin, Alice O.
REGISTRATION NUMBER: 35,601
REFERENCE/DOCKET NUMBER: 2545/90
                                                                                          Sequence 19, Application US/09385740B
Patent No. 6348320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 66.7
Matches 6, Conservative
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ORGANISM: Homo sapiens
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LOCATION: (1)..(20)

// CAPERION: Syn-corr. to C-terminal telopeptide seq. of hu. type II collagen

US-09-385-740B-18
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    Score 27; DB 4; Length 19;
Pred. No. 31;
2; Mismatches 1; Indels
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Pred. No. 31;
1; Mismatches 3; Indels
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US-09-385-740B-18
Sequence 18, Application US/09385740B
Sequence 18, Application US/09385740B
GENERAL NO. 6348120
GENERAL HYCRNATION:
APPLICANT: Eyre, David
TITLE OF INVENTION: CARTILAGE RESORPTION ASSAYS
FILE REFERENCE: WROS-1-14269
CURRENT FILING DATE: 1999-08-30
PRIOR FILING DATE: 1999-07-02
PRIOR APPLICATION NUMBER: US 60/142,274
PRIOR APPLICATION NUMBER: US 60/141,574
PRIOR PILING DATE: 1999-06-29
PRIOR FILING DATE: 1999-06-17
PRIOR FILING DATE: 1999-06-17
PRIOR FILING DATE: 1999-06-17
PRIOR FILING DATE: 1999-06-17
SPRIOR FILING DATE: 1999-06-17
      54.0%;
62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 55.6%;
Matches 5; Conservative
Query Match
Best Local Similarity 62.5
Matches 5; Conservative
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Best Local Similarity 66.7
Matches 6; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GSGFPETOP 10
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                                                                                  3 SFAKLOPR 10
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12 SFAVLEPK 19
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SAFAGLGPR 12
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Gaps

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2; Indels

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                                                                                                                                        ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-548-974-15
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                                                                                                                                                                                                                                                                               COMPUTER: IMP COMPARTIBLE
COMPUTER: IMP COMPARTIBLE
OPERATING SYSTEM: PC-DOS/MS-OOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/548,974
FILING DATE: 27-OCT-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/202,033
FILING DATE: 23-FEB-1994
ATTORNEY/AGENT INFORMATION:
NAME: Martin, Alice O.
REGISTRATION NUMBER: 35.601
REFERENCE/DOCKET NUMBER: 2545/90
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                     Matches
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                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (312) 321-4299
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
                                                                                                                                                                                                                                                 TELEFAX: (312) 321-4299 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: POTEMBA, LAN TITLE OF INVENTION: MET TITLE OF INVENTION: PC NUMBER OF SEQUENCES: 24 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 60610
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Local Similarity 40.0%;
les 4; Conservative
                                                                 Local Similarity
nes 4; Conserv
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: BRINKS HOFER GILSON & LIONE
STREET: P.O. Box 10395
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                                 1 GSSFAKLQPR 10
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GTVFSRMPPR 12
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                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lawrence A Methods And Kits For Stimulating Methods And Kits For Stimulating Production Of Megakaryocytes And Thrombocytes
                                                                                    54.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24:
                                                                                    Score 27;
Pred. No.
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Pred. No.
                                                                     Mismatches
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                                                                                      41;
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37;
                                                                                                     2
                                                                                                   Length 25;
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                                                                   Indels
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                                                                 Gaps
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RESULT 14

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RESULT 15
US-08-287-957-107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION OF TELECOMMUNICATION:

ATTORIES! AGENT INFORMATION:

NAME: Clark, Paul T. and Tsao, Y. Rocky

NAME: Clark, Paul T. and 34,053

RECISTRATION NUMBER: 30,162 and 34,053

REFERENCE DOCKET NUMBER: 00537/068001

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                              Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2
Patent No.
                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 50.0
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: HYUK K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 542-890
TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US FILING DATE: 19920727 CLASSIFICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                             APPLICANT: HYUK KIM, SUN
APPLICANT: RILEY KEYES, SUSAN
APPLICANT: MOREAU SVINTER
                                              APPLICANT: XIN DONG, ZHENG
APPLICANT: TAYLOR, JOHN
TITLE OF INVENTION: THERAPEUTIC PEPTIDE DERIVATIVES
NUMBER OF SEQUENCES: 116
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: The sequence has an amide C-terminus (i.e., COyNH2), rather the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: MS-D
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
ADDRESSEE: Fish & R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Vijayaragh APPLICANT: Archibond,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: AMINO ACID STRANDEDNESS: N/A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: N/A
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8 GTVLAKMYPR 17
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                                                                                                                                                                                                                                 55525
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                                                                                                                                                                                                                                                Application US/08287957
Massachusetts
                               225 Franklin Street
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Nagalla, Srinivasa R.
Vijayaraghavan, Srinivasan
Archibond, Anthony
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM PS/2 Model 50Z or 55SX
SYSTEM: MS-DOS (Version 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          METHOD OF CONTROLLING FERTILIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                              54.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                              2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 27; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                               DΒ
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 27;
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Mon Jan 13 09:26:19 2003
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COUNTRY: U.S.A.

ZIP: 02110-2804

ZIP: 02110-2804

COMPUTER READABLE PORM:

MEDIUM TYPE: J.* Diskette, 1.44 Mb

COMPUTER: 18 P$/2 Model 502 or 555X

COMPUTER: SIP P$/2 Model 502 or 555X

COMPUTER: Na-DOS (Version 5.0)

SOFWARRE: WordPerfect (Version 5.1)

SOFWARRE: WordPerfect (Version 5.1)

SOFWARRE: WordPerfect (Version 5.1)

APPLICATION NUMBER: US/08/287,957

FILING DATE: 09-AUG-1994

ATYORNEY/AGENT INPORMATION: 100-101

FELECOMMUNICATION INPORMATION: 100-107

FELERAX: (617) 542-8906

FEL
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Searched:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence:
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A Geneseq 101002:*

// SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
// SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Neuropeptide for c	Ghrelin-like growt	Ghrelin-like growt	Heavy chain CDR3 r	Ghrelin-like growt	Ferritin motif #16	PKB substrate #19.	Ghrelin-like growt	Ghrelin-like growt	Ghrelin-like growt	Description

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(KANG/) KANGAWA K.	26-APR-2000; 2000JP-0126623.		23-JUL-1999; 99JP-0210002.	24-JUL-2000; 2000WO-JP04907.		01-888-2001	WO200107475-A1.	Uncoinynchus mykiss.	Anguilla japonica.	Gallus gallus.	Bos taurus.	Sus scrofa.	Homo sapiens.	Rattus norvegicus.		growth hormone deficiency.	calcium concentration elevation; infant growth disorder;	Growth hormone secretagogue; GHS; ghrelin; core region;	,	Ghrelin-like growth hormone secretagogue (GHS) core region, SEQ ID NO:8.	24-WEK-2001 (III8C BUCIA)		AAB60512;	•	AAB60512 standard; peptide; 4 AA.	0512	LT 1

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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concentration in cells. The peptides are ghrelin homologues and are characterised in that a least one amino acid has been substituted by a less encompasses the unmodified peptides; the DNA encoding the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptides. The peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with common accompanying side effects. The present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence.
                                                                                 compounds which induce growth hormone secretion and calcium concentrations, useful in treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ghrelin-like growth hormone secretagogue (GHS) core region peptide #1.
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                    Matsuo H, Minamitake Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 20; DB 22; Length 4 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Growth hormone secretagogue; GHS; ghrelin; core region; calcium concentration elevation; infant growth disorder; growth hormone deficiency.
                                                                                                                                                 Claim 3; Page 184; 210pp; Japanese.
                   Новода Н,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB60535 standard; peptide; 5 AA.
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26-APR-2000; 2000JP-0126623.
                                                                                                                of infant growth disorders
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                   Kojima M,
                                                WPI; 2001-159704/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 AA;
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Gallus gallus.
Anguilla japoni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200107475-A1
                                                                               New peptide of elevate cell
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                 Kangawa K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GSSF 4
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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concratation in cells. The peptides are ghrelin homologues and are concratised in that at least one amino acid has been substituted by also encompasses the unmodified peptides; the DNA encoding the peptides, cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide. The peptides of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases comprising they are useful for treating and/or diagnosing diseases concreted by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with compounds of the invention are safe with the present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence.
                                                                                                                            New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ghrelin-like growth hormone secretagogue (GHS) core region peptide #2.
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          Minamitake Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
100.0%; Score 20; DB 22;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 4; Conservative 0; Mismatches 0;
       Matsuo H,
                                                                                                                                                                                                                                                                      Disclosure; Page 7; 210pp; Japanese.
   Hosoda H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB60536 standard; peptide; 6 AA.
                                                                                                                                                                                                          infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JUL-1999; 99JP-0210002.
29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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Kojima M,
                                                                   WPI; 2001-159704/16.
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Gallus gallus.
Anguilla japoni
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Sus scrofa.
Kangawa K,
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The period of the invention are safe with the present sequence represents a consequence.
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                                                                                                                                                              18-JUL-1996;
20-DEC-1995;
16-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                    Protein kinase B; PKB; substrate; glycogen metabolism; pancreatic cancer; regulator; protein synthesis; enzyme modulator; type II diabetes; insulin-stimulated crosstide kinase; breast cancer; ovarian cancer;
Use of protein kinase B for regulation of glycogen metabolism and protein synthesis - also peptide substrates for PKB and methods for
                                                                                                                 (MEDI-)
                                                                                                                                                                                                                                                                                                                                                      therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptide compounds which induce growth hormone secretion elevate cell calcium concentrations, useful in treatment and
                                                 WPI; 1997-341435/31.
                                                                                                                                                                                                                             20-DEC-1996;
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                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                        PKB substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 7; 210pp; Japanese
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95GB-0026083.
96GB-0010272.
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Pred. No. 7.8e+05;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Functional surrogate; analyte; affinity receptor; immunoreactive group; mimic; homogenous immunoassay; detection; diagnostic analyte; Chlamydia; cardiac marker; tumour marker; allergen; hormone; fertility; myoglobin; pregnancy; infectious disease; ferritin; myosin light chain; troponin; follicle stimulating hormone; human; growth hormone; immunoglobulin E; prolactin; parathyroid hormone; placental lactogen; hepatitis antigen; antibody; chorionic gonadotropin; luteinising hormone; cytomegalovirus; Streptcococcus; rubella; troxoplasma; DK-MB; prostate-specific antigen; carcinoembryonic antigen; alpha-fetoprotein; herpes virus; CA125.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           equivalents for regulating glycogen metabolism and/or protein synthesis is the subject of the invention. This sequence can also be used in a method of the invention for identifying agents that modulate the activity of PKB. It can also be used to screen for modulators of enzymes that catalyse PKB phosphorylation. PKB (an insulin-stimulated crosstide kinase) and its analogues etc. are used to treat disease characterised by abnormal glycogen metabolism and/or protein synthesis, especially type II diabetes and cancer (specifically of breast, pancreas and ovary). The various screening methods are used to identify agents potentially useful for treating these diseases.
This sequence represents a peptide motif derived from ferritin which important for selective binding affinity. Peptides containing motifs such as this may be used as functional surrogates in the conjugate of
                                                                                                    Labelled functional surrogate of an analyte - useful as competitor molecule in affinity assays, esp. for detecting large macromolecules such as ferritin
                                                                                                                                                                       WPI; 1997-077284/07
                                                                                                                                                                                                       Carter JM,
                                                                                                                                                                                                                                                                          07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                           W09641172-A1
                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW10773 standard; peptide; 7 AA.
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llarity 100.0%;
Conservative 0;
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                                                                 55; 156pp;
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Pred. No. 7.8e+05;
Mismatches 0;
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New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis

Disclosure; Page 180; 210pp; Japanese.

of infant growth disorders -

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the invention. The novel labelled conjugate comprises at least one label attached to a functional surrogate of an analyte of interest.

C The surrogate is capable of competing effectively with the analyte of or a limiting amount of an affinity receptor for the analyte. The conjugate exhibits an activity that is altered upon interaction with the affinity receptor and this activity can be measured and related to the analyte present in a sample. Functional surrogates compete effectively and with the analyte for a limiting amount of its such as this have an immunoreactive group that allows the surrogate to such as this have an immunoreactive group that allows the surrogate to affinity receptor. Functional surrogates are able to mimic naturally corruring analytes. They can be labelled for use in standard competitive affinity assays (sp. homogenous immunoassays) for detecting large maromolecules such as polypeptides, polysaccharides, polynuclectides, clycoproteins and lipid-containing macromolecules, as well as small haptens. Typical diagnostic analytes for detecting include cardiac or analytes associated with infectious disease. In particular, the assays create useful for detecting ferritin, parathyroid hormone, human considered with increasing hormone, numan conting macromic gonadotropin, human luteinising hormone, yethong cannot strus, DK-MB, myosin light chain, troponin, carcinoembryonic antigen, way calpha-fetoprotein, prostate-specific antigen and califormarker).
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Ghrelin-like growth hormone secretagogue (GHS) core region, SEQ ID NO:1. AAB60507 standard; peptide; 7 AA. 24-APR-2001 (first entry) 1 GSSF 4 3 GSSF 6 AAB60507; **AAB**60507 셤 ઠે

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Growth hormone secretagogue; GHS; ghrelin; core region; calcium concentration elevation; infant growth disorder; growth hormone deficiency. 24-JUL-2000; 2000WO-JP04907 Bos taurus. Gallus gallus. Anguilla japonica. Oncorhynchus mykiss. Rattus norvegicus, WO200107475-A1. Sus scrofa. 01-FEB-2001.

The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion cludes the secretion of growth hormone and/or elevates calcium ion cludes to concentration in cells. The peptides are ghrelin homologues and are caracterised in that at least one amino acid has been substituted by also encompasses the unmodified peptides; the DNA encoding the peptides comprising recombinant production, optionally followed by the invention, and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases the invention are useful for treating and/or diagnosing diseases particular, they are useful for treating and/or diagnosing diseases particular, they are useful for promoting infant growth due to growth connent deficiency. The compounds of the invention are safe with common deficiency. The compounds of the invention are safe with a accompanying side effects. The present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence. ô antibody; monoclonal; chimaeric; grafted; humanised; IL-2; interleukin-2; cytokines; interleukin-2 receptor; 55kD beta chain; activated T cells; T cell mediated disease; graft versus host disease; transplant rejection; autoimmune diseases; chemotherapy; immunosuppressants; T cell typing; diagnosis; testing; detection; ss. Gaps Heavy chain CDR3 region for humanised anti-IL-2 receptor antibody. Human interleukin-2 receptor antibodies - useful for treating and preventing T-cell mediated diseases e.g. graft versus host disease, transplant rejection etc. .. o Length 7; 0; Indels 100.0%; Score 20; DB 22; 100.0%; Pred. No. 7.8e+05; iive 0; Mismatches 0; Winter GP; AAR32118 standard; Protein; 8 AA. Waldmann H, Query Match Best Local Similarity 100.00 These 4; Conservative 92WO-GB01258. 91GB-0015010. (WALD/) WALDMANN H. (WELL) WELLCOME FOUND LTD. 28-MAY-1993 (first entry) Crowe JS, Lewis AP, WPI; 1993-045493/05. N-PSDB; AAQ36579 Rattus rattus 10-JUL-1992; 11-JUL-1991; 1 GSSF 4 21-JAN-1993. 1 GSSF 4 WO9301289-A. Sequence AAR32118;

Hosoda H, Matsuo H, Minamitake Y;

Kangawa K, Kojima M,

(KANG/) KANGAWA K.

WPI; 2001-159704/16.

29-NOV-1999; 99JP-0338841. 26-APR-2000; 2000JP-0126623.

23-JUL-1999;

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Matches
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        The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptide also encompasses the unmodified peptides; the DNA encoding the peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Growth hormone secretagogue; GHS; ghrelin; core region; calcium concentration elevation; infant growth disorder;
                                                                                                                                                                   New peptide compounds which induce growth hormone secretion elevate cell calcium concentrations, useful in treatment an
                                                                                                                                                                                                                                                                                                           23-JUL-1999; 99JP-0210002
29-NOV-1999; 99JP-0338841
26-APR-2000; 2000JP-0126623
                                                                                                                       Disclosure; Page 7; 210pp; Japanese
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                                                                                                                                                                                                                                                                               (KANG/) KANGAWA K.
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                                                                                                                                                    growth disorders
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                                                                                                                                                                                                                                                 Kojima
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larity 100.0%;
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99JP-0338841.
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Pred. No. 7.8e+05;
Mismatches 0;
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 Buch
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 DNA; a
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the peptides; oducing the
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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an annibody specific for a peptide. The peptides invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptid of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence.
                                                                                                                                                                                                    New peptide compounds which induce gelevate cell calcium concentrations, of infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 growth hormone
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                                                                                                                                                                       Disclosure; Page 7;
                                                                                                                                                                                                                                                                                     Kangawa
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29-NOV-1999;
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scrofa.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hormone secretagogue; GHS;
concentration elevation;
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; 99JP-0338841.
; 2000JP-0126623.
                                                                                                                                                                                                                                                                                       Kojima M,
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Pred. No.
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                                                                                                                                                                                                                 rowth hormone secretion useful in treatment and
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cheв 0;
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diagnosis
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A set of peptide ligands consisting of specific complementary peptides to proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided in the specification.
                              Human; complementary peptide; ligand; drug discovery; drug design.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Page 344; 646pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 drug candidates or pro-drugs
                                                                                                                                                                                                                                              99GB-0029464.
                                                                                                                                                                                                    13-DEC-2000; 2000WO-GB04776.
                                                                                                                                                                                                                                                                                                                                 Heal JR;
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Best Local Similarity
4; Conserv?
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                                                                          Homo sapiens
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caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The peptide, which can be isolated from the ganglion of Helix pomatia or can be synthesised, has the activity to improve muscular contraction, as demonstrated by tests using anterior byssus retractor
                                                                                                                                                                                                                        Gaps
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100.0%; Score 20; DB 22;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 4; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 1.3e+02;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neuropeptide for controlling muscle contraction.
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                                                                                                                               9 AA;
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    Length 10;
                                              Indels
100.0%; Score 20; DB 22;
ilarity 100.0%; Pred. No. 1.3e+02;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                           Human complementary peptide, SEQ ID NO: 2137.
                                                                                                                                                                                                                                   AAG95943 standard; Peptide; 10 AA.
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RESULT 13
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                                                                                                  generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as respents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided in the specification.
                                                                                                                                                                                                                                     A set of peptide ligands consisting of specific complementary peptides to proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are
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                                                                               Sequence
                                                                                                                                                                      The invention relates to a set of complementary peptide ligands
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RESULT 15
AAG85523
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AC AAG85
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AAG95981
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                                            Saccharomyces cerevisiae; drug discovery; drug desi
                                                                              Saccharomyces cerevisiae peptide,
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                                                                                                                            AAG85523;
                                                                                                                                                   AAG85523 standard; Peptide; 10
                                                                                                                                                                                                                                                                                                     Sequence
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                                           drug design
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WO200142276-A1

Saccharomyces cerevisiae

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The invention relates to the identification of complementary peptides by analysis of protein and nucleotide sequence databases from higher eukaryotic genomes, excluding human and plants. The specific complementary peptides interact with their relevant target proteins encoded in the eukaryote genome. The peptides may be used as reagents and drugs for drug discovery and as lead ligands for drug design and development. The present sequence is a complementary peptide from Saccharomyces cerevisiae.
                                                                                                                                                                                                                         Identifying complementary peptides by analysis of protein and nucleotide sequence databases, useful in drug design -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
100.0%; Score 20; DB 22; Length 10;
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 4; Conservative 0; Mismatches 0; Indel8
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                                               13-DEC-2000; 2000WO-GB04773.
                                                                                     99GB-0029471
                                                                                                                                                         Roberts GW, Heal JR;
                                                                                                                      (PROT-) PROTEOM LTD.
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Qy 1 (Db 28 (Query Match Best Local Matches	RESULT 2 PH1738 Ig heavy chain V regicly species: Mus muscul C; Species: Mus muscul C; Cpate: 24-Feb-1994 # C; Accession: PH1738 R; McHeyzer-Williams; J Exp. Med. 178. 295 A; Title: Antigen-driv A; Reference number: PA; Accession: PH1738 A; Molecule type: mRNA A; Residues: 1-33 cMCH A; Experimental source C; Superfamily: immuno C; Keywords: heterotet	Query Matcl Best Local Matches Qy 1 GSI Db 20 GSI	R;MCHeyzer-Williams, t J: Exp. Med. 178, 292 A;Title: Antigen-drive A;Reference number: PA A;Recession: PH.733 A;Molecule type: mRNA A;Residues: 1-25 <mch: A;Experimental source C;Superfamily: immunoc C;Keywords: heteroteti</mch: 	RESULT 1 PH1733 I PH273 C/Species C/Species C/Date: 24 C/Accession	44444333333333333333333333333333333333
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A;Cross-references: EMBL:AF020713; NID:g3025478; PID:g3025626; PIDN:AAC13121.1

E;Kunst, F; Ogasawara, N; Moszer, I.; Albertini, A.M.; Alloni, G; Azevedo, V.; Berres, Denn, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Ch. A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E. Akuthors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F. Akuthors: Lauber, J.; Lazarevil, G.; Kwogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois A; Authors: Lauber, J.; Lazarevil, V.; Persk, S.H.; Parro, V.; Pohl, T.M.; Portetell, M.; M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetell, A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seronakeuchi, M.; Tamakoshi, A.; Tanaka, T.; Targetra, P.; Tosato, V.; Uchiyama, T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, A; Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis. A; Reference number: A69580; MUID:98044033; PMID:9384377
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A;Title: Basic fibroblast growth factor induces 3T3 fibroblasts to synthesize and secret A;Reference number: A56814; MUID:92031730; PMID:1932134
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 24-Feb-13-4-Feb-1994 #text_change 17-Mar-1999
C;Accession: PH1739; PH1737
R;MCHQYZer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
J. Exp. Med. 178, 255-307, 1993
A;Fitle: Antigen-driven B cell differentiation in vivo.
A;Reference number: PH1675; MUID:93301607; PMID:8315385
Ig heavy chain V region (clone NP-12-8) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 17-Mar-1999
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                                         Cispecies: Mus musculus (house mouse)
Cipate: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 17-Me CiAccession: PHI/Jep.
CiAccession: PHI/Jep.
J. Exp. Med. 178, 295-307, 1993
J. Exp. Med. 178, 295-307, 1993
J. Fitle: Antigen-driven B cell differentiation in vivo.
A;Reference number: PHI675; MUID:93301607; PMID:8315385
A;Recession: PHI742
A;Molecule type: mRNA
A;Residues: 1-33 400H)
A;Residues: 1-33 400H)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A, Experimental source: B cell, clone NP-12-3
C, Superfamily: immunoglobulin V region; immunoglobulin homology
C, Keywords: heterotetramer; immunoglobulin
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A,Residues: 1-33 <MCH>
A,Experimental source: B cell, clone NP-12-5
A,Accession: PH1737
A,Molecule type: mRNA
A,Residues: 1-33 <MCH2>
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A; Residues: 1-38 < DAV>
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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A; Accession: PH1739
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rap60B protein - Bacillus subtilis plasmid pTA1040
C;Species: Bacillus subtilis
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 15-Oct-1999
C;Cacession: 140555
R;Meijer, W.J.; Venema, G.; Bron, S.
R)Woleic Acids Res. 23, 612-619, 1995
A)Title: Characterization of single strand origins of cryptic rolling-circle plasmids f
A;Reference number: 140549; MUID:95206941; PMID:7899081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Species: Bacillus subtilis phage SPBC2
C; Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 15-Oct-1999
C; Accession: T12912; C69925
R; Lazarevic, V.; Duesterhoeft, A.; Soldo, B.; Hilbert, H.; Mauel, C.; Karamata, D. submitted to the EMBL Data Library, August 1997
A; Description: The complete nucleotide sequence of the Bacillus subtilis SPbetac2 proph. A; Reference number: 217583
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A;Experimental source: plasmid pTA1040
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                                                                                                              100.0%; Score 20; DB 2; Length 38; larity 100.0%; Pred. No. 95; Conservative 0; Mismatches 0; Indels
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98;
A;Note: sequence modified after extraction from NCBI backbone C;Superfamily: peptidylprolyl isomerase; cyclophilin homology C;Keywords: cis-trans-isomerase
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A;Molecule type: DNA
A;Residues: 1-39 <RES>
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A;Molecule type: DNA
A;Residues: 1-39 <LAZ>
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larity 100.0%; Pred. No. 98;
Conservative 0; Mismatches
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Best Local Similarity
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                                                                                                              Query Match
Best Local Similarity
Matches 4; Conserv
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R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; DeHoff, B. e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, Y. P.; Sun, P.M.; Minkler, M.E. J. Bacteriol. 183, 5709-5717, 2001
J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas, A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A;Reference number: A97872; MUID:21429245; PMID:11544234
A;Accession: B97874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, 1 R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-49 < KUR>
                                                                                                                                                                                                                                                                                                                                                                                               degenerate transposase (orf2) [imported] - Streptococcus pneumoniae (strain C;Species: Streptococcus pneumoniae C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 22-Oct-2001 C;Accession: B97874
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A; Residues: 1-42 <HEI>
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A/Status: nucleic acid
A/Molecule type: DNA
A/Residues: 1-39 <KUN>
                                                                                                            A;Gene: IS1167-truncation
                                                                                                                                                                                                               A;Status: preliminary
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Query Match
Best Local Similarity 100.
                                                                                                                                                  Cross-references: GB:AE007317; PIDN:AAK98822.1; PID:g15457547; GSPDB:GN00174
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Ol; strain N16961; biotype El Tor
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Pred. No.
                           Score 20; DB 2;
Pred. No. 1.2e+02;
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I.; Sellers,
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R,Johnsen, A.H.; Rehfeld, J.F.
Bur. J. Biochem. 213, 875-879, 1993
A;Title: LymnaDFamides, a new family of neuropeptides from the A;Reference number: S32471; MUID:93238777; PMID:8477756
A;Accession: S32473
                                                                                                                                                                                      rymnaDFamide 3 - great pond snail
C;Species: Lymnaea stagnalis (great pond snail)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
C;Accession: S32473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: EMBL:X67386; NID:g50927; PIDN:CAA47798.1; PID:g1333920 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
A;Cross-references: PIDN:AAB26364.1; PID:g299831
A;Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
                                                          A; Molecule type: protein A; Residues: 1-13 < JOH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             coagulation factor X inhibitor - sharp-nosed viper (fragment)
C;Species: Agkistrodon acutus (sharp-nosed viper)
C;Date: 03-May-1994 #sequence_revision 03-May-1994 #text_change 03-May-1994
C;Accession: A49261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
S25056
                                                                                                                                                                                                                                                                      $32473
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A; Accession: A49261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                submitted to the EMBL Data Library, July 1992 A;Description: In situ studies on the primary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Mus musculus (house mouse)
C;Date: 25-Feb-1994 #sequence_revision 01-Dec-1995 #text_change 20-Jun-2000
                                                                                                                                                                                                                                                                                            RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Toxicon 31, 1445-1457, 1993
A; Title: Coagulation factor
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A; Residues: 1-12 < JAC>
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A; Accession: S25056
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Cispecies: Homo sapiens (man)
Cispecies: Homo sapiens (man)
Cispecies: Homo sapiens (man)
Cispecies: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
Ciscossion: 566627
RiMoreno, F.J.; Medina, M.; Perez, M.; Montejo de Garcini, E.; Avila, J.
RiMoreno, F.J.; Medina, M.; Perez, M.; Montejo de Garcini, E.; Avila, J.
RiMoreno, F.J.; Medina, M.; Perez, M.; Montejo de Garcini, E.; Avila, J.
RiMorence number: 566627; MulD:96032547; PMID:7556645
A;Accession: 56627
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C;Species: Lotus corniculatus
C;Date: 15-Jun-1990 #sequence_revision 15-Jun-1990 #text_change 04-Mar-2000
C;Date: 15-Jun-1990 #sequence_revision 15-Jun-1990 #text_change 04-Mar-2000
C;Date: 15-Jun-1990 #sequence_revision 15-Jun-1990 #text_change 04-Mar-2000
R;Accession: B2711
R;County and J; Petersen, T.E.; Marcker, K.A.
Proc. Natl. Acad. Sci. U.S.A. 84, 5754-5757, 1987
Proc. Natl. Acad. Sci. U.S.A. 84, 5754-5757, 1987
A;Title: Expression of a complete soybean leghemoglobin gene in root nodules of transgen A;Accession: B32711
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C;Species: Mus sp. (mouse)
C;Accesion: 167525
R;Chies, J.A.; Lembezat, M.P.; Freitas, A.A.
C;Accesion: 167525
R;Chies, J.A.; Lembezat, M.P.; Freitas, A.A.
A;Title: Entry of B lymphocytes into the persistent cell pool in non-immunized mice is 1A;Reference number: 153392; MUID:9429870; PMID:8026526
A;Accession: 167525
A;Accession: 167525
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-15 cRES>
A;Residues: 1-15 cRES>
A;Cross-references: GB:S71349; NID:9550037
C;Genetics:
A;Gene: Ig VH7183
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F;13/Modified site: amidated carboxyl end (Phe) #status predicted
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85.0%; Score 17; DB 2; Length 15;
Best Local Similarity 75.0%; Pred. No. 2.5e+02;
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Pred. No. 3.7e+02;
1; Mismatches 0; Indele
A;Molecule type: protein
A;Reaidues: 1-22 <STO>
Superfamily: globin; globin homology
C;Keywords: oxygen carrier
                                                                                         vuery Match
Best Local Similarity 75.0%;
Matches 3; Conservative
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 10, 2003, 15:55:16; Search time 3.63636 Seconds (without alignments)
45.624 Million cell updates/sec

Title: A
Perfect score: 20
Sequence: 1 gssf 4
Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
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Minimum DB seq length: 0
Maximum DB seq length: 50
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Total number of hits satisfying chosen parameters:

3754

112892 seqs, 41476328 residues

Searched:

Database : SwissProt_40:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

CERB_CERCA CERB_CERCA ID	RX MEDLINE=93238777; PubMed=8477756; RA Johnsen A.H., Rehfeld J.F.; RT "LymnaDFamides, a new family of neuropeptides from the pond snail, RT LymnaDeanides, a new family of neuropeptides from the pond snail, RT invertebrates?"; RL Eur. J. Biochem. 213:875-879(1993). CC -! -SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY. DR PIR; \$32473; \$32473. KW Neuropeptide; Amidation. FT MOD RES 13 13 AMIDATION. FT UNSURE 12 12 SQ SEQUENCE 13 AA; 1462 MW; 9CA07BA3F5D5B865 CRC64; Query Match 85.0%; Score 17; DB 1; Length 13; Best Local Similarity 75.0%; Pred. No. 88; Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps Qy 1 GSSF 4 USSF 4 USSF 4 USSF 15 10	RESULT 1 NP3_LYMST ID
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WA MEDLINE=9347989; PubMed=8346031;
WA Hallock R.B., Hong L., Stutez E.;
WA Hallock R.B., Hong L., Stutez E.;
A Orsat B., Spielmann A., Stutez E.;

R. Complete sequence of Euglena gracilis chloroplast DNA.";
R. Nucleic Acids Res. 21:337-354(1993).
C CROSS-MEMBRANE TRANSPORT OF ELECTRONS BETWEEN PHOTOSYSTEM II AND CC I, AS WELL AS IN CYCLIC ELECTRON BLOW AROUND PHOTOSYSTEM II.
PETG IS REQUIRED FOR EITHER THE STABILLTY OR ASSEMBLY OF THE CYTOCHROME B6-F COMPLEX.
C CYTOCHROME B6-F COMPLEX.
C CYTOCHROME B6-F COMPLEX.
C C-:-SIMILARITY: BELONGS TO THE PETG FAMILY.
"Purification and primary structure of ceratotoxin A and B, two antibacterial peptides from the female reproductive accessory glands of the medfly Ceratitis capitata (Insecta:Diptera)."; Insect Biochem. Mol. Biol. 23:591-598(1993)."; GRAM-POSITIVE AND GRAM-NEGATIVE BACTERIA. THEY HAVE AS WELL HEMOLYTIC ACTIVITY. THESE PROTEINS ARE STABLE EVEN AT 100 DEGREES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-APR-1993 (Rel. 25, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Cytochrome B6-F complex subunit V (Cytochrome b6f complex subunit
                                                                                                                                                                                                                                              -!- SUBUNIT: HOMOPOLYMER OF FOUR TO SIX SUBUNITS.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: STRUCTURALLY RELATED TO CECROPINS, DEFENSINS AND
APIADECHINS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 17; DB 1; Length 29;
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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Chloroplast.
Eukaryota; Euglenozoa; Euglenida; Euglenales; Euglena.
NCBI_TaxID=3039;
                                                                                                                                                                                                                                                                                                                                                                       Insect immunity, Hemolysis, Antibiotic.
SEQUENCE 29 AA, 2861 MW; BE57F4EECB2DA6B0 CRC64;
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STROMAL (POTENTIAL).
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01-APR-1993 (Rel. 25, Last seq
16-OCT-2001 (Rel. 40, Last ann
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P30396;
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(Rel. 14, Last sequence update)
(Rel. 41, Last annotation update)
(Gamma-carboxyglutamic acid-containing protein) (Bone Gla-
                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                        Bacteriophage T7.
Viruses; dsDNA viruses, no RNA stage; Caudovirales; Podoviridae;
T77-like viruses.
NCBI_TaxID=10760;
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                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=83241725; PubMed=6864790;
Dunn J.J., Studier F.W.;
"Complete nucleotide sequence of bacteriophage T7 DNA and the
                       Score 17; DB 1; Length 37;
Pred. No. 2.6e+02;
1; Mismatches 0; Indels
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13806339E110D3D6 CRC64;
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Pred. No. 3.4e+02;
1; Mismatches 0;
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21-JUL-1986 (Rel. 01, Last sequence update)
01-WAR-1989 (Rel. 10, Last annotation update)
Gene 0.5 protein.
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                                                                                                                                                                                                                                                                                                                                                        locations of T7 genetic elements.";
J. Mol. Biol. 166:477-535(1983).
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EMBL; V01127; CAA24329.1; -.
PIR; A04402; W0BP57.
SEQUENCE 47 AA; 4745 MW;
                        85.0%;
75.0%;
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75.0%;
4147 MW;
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37 AA;
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17 GASF 2
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P03777;
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DT 01-APR.
DT 15-JUN.
DE OSTCOCC.
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01-FEB-1994 (Rel. 2

16-OCT-2001 (Rel. 4
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SEQUENCE
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InterPro; IPR002294; VitK_dep_GLA.
Pfam; PF00594; gla; 1
PRINTS; PR00002; GLABONE.
SMART; SM00069; GLA; 1.
PROSITE; PS00011; GLU_CARBOXYLATIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "The amino acid sequence of Emu osteocalcin: gas phase sequencing Gla-containing proteins";
Biochem. Int. 15:271-277 (1987).
-I- FUNCTION: CONSTITUTES 1-2% OF THE TOTAL BONE PROTEIN. IT BINDS STRONGLY TO APATITE AND CALCIUM.
-I- PTM: GAMMA-CARBOXYGLUTAMIC ACID RESIDUES ARE FORMED BY VITAMIN DEPENDENT CARBOXYLATION. THESE RESIDUES ARE ESSENTIAL FOR THE
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                                                                                                                          MEDLINE=92210010; PubMed=1555775; Chavrier P., Simons K., Zerial M.; The complexity of the Rab and Rho GTP-binding revealed by a PCR cloning approach."; Gene 112:261-264(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dromaius novae-hollandiae (Emu).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                     Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                 Ras-related protein RAB22A OR RAB22.
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                                                                                                                                                                                            TISSUE=Kidney;
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SIMILARITY: BELONGS TO THE OSTEOCALCIN / MATRIX GLA-PROTEIN
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40, Last annotation updat
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Rodentia;
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75.0%;
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Sciurognathi;
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3.5e+02;
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thi; Muridae; Murinae; Mus
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RESULT 8
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P08766;
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01-FEB-1991 (Rel. 17, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Leucokinin V (L-V).
Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria;
Blaberoidea; Blaberidae; Leucophaea.
                                                         01-NOV-1988 (Rel. 09,
01-NOV-1988 (Rel. 09,
01-NOV-1997 (Rel. 35,
Small core protein (J
                                                                                                                                                                                                                                                                                                    Holman G.M., Cook B.J., Nachman R.J.;
"Isolation, primary structure, and synthesis
myotropic peptides of Leucophaea maderae.";
Comp. Biochem. Physiol. 88C:27-30(1987)
-!- FUNCTION: THIS CEPHALOMYOTROPIC PEPTIDE S
ACTIVITY OF COCKROACH PROTODEUM (HINDGUT)
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MGD; MGI:105072; Rab22.
     Bacteriophage alpha-3, a
Bacteriophage phi-K.
Viruses; ssDNA viruses;
VIRI TaxID=10849, 10848;
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nilarity 75.0%;
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                  Microviridae; Microvirus
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736365A5B9C865B8 CRC64;
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Pred. No. 1.1e+05
0; Mismatches
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SIMILARITY).
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                                                                                                                                                                                                                                                                                                                                                 [3]
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
SPECIES=Phage phi-K;
Kodaira K.-I., Tadokoro H., Suzuki K., Oki M., Manto S., Taketo A.;
Kodaira K.-I., Tadokoro H., Suzuki K., Oki M., Manto S., Taketo A.;
Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
I- FUNCTION: THE J PROTEIN IS ASSOCIATED WITH THE DNA AND IS SITUATED IN AN INTERIOR CLEFT OF THE F PROTEIN.
IN AN INTERIOR CLEFT OF THE F PROTEIN.
J PROTEINS, AND 12 COPIES OF THE H PROTEIN.
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"Isolation and characterization of a novel 39 kilodalton whey protein
from bovine mammary secretions collected during the nonlactating
                  SPECIES=Phage alpha-3;
MEDLINE=92223109; PubMed=1532908;
Kodaira K.-I., Nakano K., Okada S., Taketo A.;
"Nuclectide sequence of the genome of the bacteriophage alpha 3:
interrelationship of the genome structure and the gene products with those of the phages, phi X.74, G4 and phi K.";
Biochim. Biophys. Acta 1130:277-288(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gарв
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Bukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidea; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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16-OCT-2001 (Rel. 40, Last annotation update)
Chitinase-3 like protein 1 (Cartilage glycoprotein-39) (GP-39)
Whey protein) (Fragment).
                                                                                                                                                                                                                                                               Kodaira K.-I., Taketo A.; "Isolation and some properties of bacteriophage alpha3 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16; DB 1; Length 24;
Pred. No. 3.1e+02;
0; Mismatches 1; Indels
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                                                                                                                                                                                             SEQUENCE FROM N.A.
SPECIES=Phage alpha-3;
MEDLINE=84294906; Pubmed=6088949;
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EMBL; X60374; CAA42849.1; -.
EMBL; X6033; CAA42890.1; -.
PIR; S09546; S09546.
PIR; S22333; S22333.
COAT protein; DNA-binding.
SEQUENCE 24 AA; 2823 MW; 01
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larity 75.0%;
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P30922;
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DT 01-JUL-
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch)
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Eukaryota; Rhodophyta; Bangiophyceae; Porphyridiales; Porphyridiaceae;
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Biochem. Biophys. Res. Commun. 150:329-334(1988).
-!- FUNCTION: MY PLAY DAN IMPORTANT ROLE IN THE CAPACITY OF CELLS RESPOND TO AND COPE WITH CHANGES IN THEIR ENVIRONMENT.
-!- SUBCELLULAR LOCATION: EXtracellular.
-!- TISSUE SPECIFICITY: MAMMARY SECRETIONS COLLECTED DURING THE NONLACTATING PERIOD.
-!- PTM: GLYCOSYLATED.
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Plant Mol. Biol. 24:253-257(1994).
-!- SIMILARITY: BELONGS TO THE L27P FAMILY OF RIBOSOWAL PROTEINS.
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PIR; A27682, A27682.
InterPro; IPR001579; Chitinase_18/2.
InterPro; IPR001223; Glyco_hydro_18.
ProDom; PD000471; Glyco_hydro_18; 1.
ProDom; PD000471; Glyco_hydro_18; 1.
PROSITE; PS01095; CHITINASE_18; PARTIAL.
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Pred. No. 4.8e+02;
0; Mismatches 1; Indels
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Pred. No. 6.4e+02;
0; Mismatches 1; Indels
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16-OCT-2001 (Rel. 40, Last annotation update)
Chloroplast 50S ribosomal protein L27 (Fragment)
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01-NOV-1995 (Rel. 32, Last seq
16-OCT-2001 (Rel. 40, Last ann
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Les 3; Conserv
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NON TER 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Netzker R., Koechel H.G., Basak N., Kuentzel H.; "Nucleotide sequence of Aspergillus nidulans mitochondrial genes coding for ATPase subunit 6, cytochrome oxidase subunit 3, seven unidentified proteins, four tRNAs and L-rRNA."; Nucleic Acids Res. 10:4783-4794(1982).
                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae; Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                             PIR; A04518; QXASF.

Hypothetical protein; Mitochondrion.

SEQUENCE 48 AA; 5628 MW; FC6AE8D1CD1992DD CRC64;
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 SEQUENCE, AND MASS SPECTROMETRY.
                           NCBI_TaxID=94770;
                                                                                                Litoria citropa (Australian blue mountains tree frog).
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Holman G.M., Cook B.J., Nachman R.J.;
"Isolation, primary structure, and synthesis of leucokinins V and myotropic peptides of Leucophaea maderae.";
Comp. Biochem. Physiol. 88C:27-30(1987).
-i- FUNCTION: THIS CEPHALOMYOTROPIC PEPTIDE STIMULATES CONTRACTILE ACTIVITY OF COCKROACH PROTODEUM (HINDGUT).
-i- SIMILARITY: TO THE OTHER LEUCOKININS, AND TO MANDUCA SEXTA ANE HELIOTHIS ZEA ADIPOKINETIC HORMONE.
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MOD_RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria;
Blaberoidea; Blaberidae; Leucophaea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-FEB-1991 (Rel. 17, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              using electrospray mass spectrometry.";
Rapid Commun. Mass Spectrom. 13:2498-2502(1999).
-i- FUNCTION: HYPOTENSIVE NEUROPEPTIDE (PROBABLE).
-i- TISSUE SPECIFICITY: SECRETED BY THE SKIN DORSAL
-i- PTM: ISOFORM 4.274 DIFFERS FROM ISOFORM 4.2 IN N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P19988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; FALSE_NEG.
Amphibian skin; Hypotensive agent; Amidation; Sulfation.
Amphibian skin; Hypotensive agent; Amidation; Sulfation.
MOD_RES 1 PYRROLIDONE CARBOXYLIC ACID
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Pred. No. 2.6e
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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Periviscerobinin-2 (Pea-PW-2).
Periviscerobinin-2 (Pea-PW-2).
Periplaneta americana (American cockroach).
Pukaryota, Metazoa, Arthropoda, Mandibulara, Pancrustacea, Hexapoda,
Insecta, Pterygota, Neoptera, Octhopteroidea, Dictyoptera, Blattaria,
Blattoidea, Blattidae, Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                          MEDIANE-20064325; PubMed=10988292;
Balaji R.A., Ohtake A., Sato K., Gopalakrishnakone P., Kini R.M., Seow K.T., Bay B.-H.;
"Lambda-conotoxins, a new family of conotoxins with unique disulfide pattern and protein folding. Isolation and characterization from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Venom of Conus marmoreus.";
J. Biol. Chem. 275:39516-39522 (2000).
-!- FUNCTION: Inhibits the neuronal noradrenaline transporter.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFOITY: Expressed by the venom duct.
-!- MASS.SPECIROITY: MW=1262.77; MW ERR=0.07; METHOD-Electrospray.
-!- SIMILABITY: BELONGS TO THE CHI/LAMBDA-CONOTOXIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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"Isolation of periviscerokinin-2 from the abdominal perisympathetic organs of the American cockroach, Periplaneta americana.";
Peptides 19:801-809(1999)
-!- FUNCTION: MYOACTIVE PEPTIDE; HAS EXCITATORY ACTIONS ON THE HYPERNEURAL MUSCLE.
-!- MASS SPECTROMETRY: MW=1189.3; METHOD=MALDI.
MOD_RES 12 AMIDATION.
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SEQUENCE, FUNCTION, AND MASS SPECTROMETRY.

MEDLINE-98326577; Pubmed-9664444;

Predel R., Rapus J., Eckert M., Holman G.M., Nachman R.J., Wang Y.,
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0
15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Lambda-conotoxin CMrX.
Conus marmoreus (Marble cone).
Eukaryota; Mctazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
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SEQUENCE 12 AA; 1190 MW; 2F4D8EEE1E805728 CRC64;
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Matches 3; Conservative
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Maximum Match 100%
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Ospn7 macaca mula
Ogfjb1 arabidopsis
O45446 bacillus su
O64160 bacteriopha
O91137 mus musculu
O31888 bacillus su
O9477 vibrio chol
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Q77894 human immun	Q77893 human immun	Q77892 human immun	Q77890 human immun		Q77901 human immun	Q77922 human immun	Q77920 human immun	Q77915 human immun	Q77918 human immun		Q77928 human immun	Q77927 human immun	Q77924 human immun	Q77926 human immun	Q77925 human immun	Q77923 human immun	Q79996 human immun	Q79997 human immun		O90520 human immun			84	O68187 lactococcus			Q900f0 human immun	Q900f1 human immun

ALIGNMENTS

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RESULT 2
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Matches 4
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Q9QVH8;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
SUPERINDUCIBLE protein 24, SIP24=CYCLOPHILIN homolog,
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            Q8SPN7
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
MCBI_TaxID=10095;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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InterPro; IPR002130; CSA_PPIase.
Pfam; PF00160; pro_isomerase; 1.
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17 AA; 1785 MW;
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            PRELIMINARY;
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            PRT;
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Bacillus subtilis.
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nes 4; Conserv
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=COLUMBIA;
MEDLINE=99087489; PubMed=9872454;
Nakamura Y., Sato S., Asamizu E., Kaneko T., Kotani H., Miyajima N.,
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DNA Res. 5:297-308 (1998).
EMBL, AB015477; BAB087071; -.
SEQUENCE 38 AA, 4298 MW; 0340720C8BFB476B CRC64;
                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae, Cercopithecinae, Macaca.
                                                                                                                                                                                                                                                                 Norgren R.B. Jr., Zink M.A., Jia Y., Ojeda S.R., Spindel E.R., "Construction of a targeted rhesus macaque microarray."; Submirted (WAR-2002) to the EMBL/GenBank/DDBJ databases. EMBL, AV083584; AMN1199.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 20; DB 6; Length 34; — 100.0%; Pred. No. 2.4e+02; tive 0; Mismatches 0; Indels 0
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                 01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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Last sequence update)
Last annotation update)
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                                                                                                                       Macaca mulatta (Rhesus macaque)
                                                                                Slit-like protein 2 (Fragment).
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Best Local Similarity
Matches 4; Conserv
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01-NOV-1996 (01-NOV-1996 (01-NOV-1998 (01-NOV-199
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Q9FJB1;
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                                                                                                                                                           Meijer W.J., Venema G., Bron S.;
"Characterization of single strand origins of cryptic rolling-circle plasmids from Bacillus subtilis";
Nucleic Acids Res. 23:612-619(1995).
EMBL; U32378; AAC44412.1; -.
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Cyclophilin A (Fragment).

Cyclophilin A (Fragment).

Mus musculus (Mouse)

Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

[1]
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NCBI_TaxID=66797;
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Plasmid pTA1040.
Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillales;
Bacillaceae; Bacillus.
NCBI_TaxID=1423;
                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 20; DB 2; Length 39; 100.0%; Pred. No. 2.7e+02; ive 0; Mismatches 0; Indels (
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Karamata D.;
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Submitted (AUG-1997) to the EMBL/GenBank/DDBJ databages.
EMBL; AF020713; AAC13121.1;
Hypothetical protein.
SEQUENCE 39 AA; 4090 MW; AC22034F15E873E1 CRC64;
                                                                                                                                                                                                                                                                                                                                                         39 AA; 4125 MW; CO86DA26EB01AE03 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-AUG-1998 (TrEMBLrel. 07, Created)
01-AUG-1998 (TrEMBLrel. 07, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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Last annotation update)
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                                                                                                                                                      STRAIN=IAM1232;
MEDLINE=95206941; PubMed=7899081;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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RESULT 7

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AC 03188

AC 0318

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                 RA Kunst F. Ogasawara N. Moscer I., Albertini A.M., Alloni G.,
RA Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
RA Borriss R., Boursier L., Brans A., Braun M., Brignell S.C., Bron S.,
RA Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
RA Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.K., Errington J., Fabret C., Ferrari E., Foulger D.,
RA Entlan K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
RA Entlan K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
RA Gimm S.Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Gimm S.Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Gimm S.Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Guiseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A.,
RA Hilbert H., Holsappel S., Hosono S., Hullo M.F., Itaya M., Jones L.,
RA Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M., Klein C.,
RA Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
RA Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
RA Medina N., Mellado R.P., Mizuno M., Mosetl D., Nakai S., Noback M.,
RA Moone D., O'Reilly M., Ogawa K., Ogiwara A., Oudega B., Park S.H.,
RA Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie Y.,
RA Sato T., Scanlan E., Schleich S., Schroeter R., Scoffone F.,
RA Sekiguchi J., Sekowska A., Seror S.J., Serror P., Shin B.S., Soldo B.,
RA Aseuchi M., Tanconi E., Takagi T., Takahashi H., Takemaru K.,
RA Yoshida K., Yoshikawa H.F., Zumstein E., Yoshikawa H., Danchin A.,
The Complete genome sequence of the gram-positive bacterium Bacillus
RT The Complete genome sequence of the gram-positive bacterium Bacillus
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EMBL; AF171073; AAD50996:1; -.
HSSP; P05092; 2CPL.
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01-JAN-1998 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
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Bacteria; Firmicutes;
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InterPro; IPR002130; CSA_PPIase.
Pfam; PF00160; pro_isomerase; 1.
PROSITE; PS50072; CSA_PPIASE_2; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20422670; PubMed=10964515;
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Pred. No. 2.
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2.7e+02;
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MEDLINE=20406833; PubMed=10952301;
Heidelberg J.F., Eisen J.A., Nelson W.C., Clayton R.A., Gwinn M.L.,
Dodson R.J., Haft D.H., Hickey E.K., Peterson J.D., Umayam L.A.,
Gill S.R., Nelson K.E., Read T.D., Tettelin H., Richardson D.,
Ermolaeva M.D., Vamathevan J., Bass S., Qin H., Dragoi I., Sellers P.,
McDonald L., Utterback T., Fleischmann R.D., Nierman W.C., White O.,
Salzberg S.L., Smith H.O., Colwell R.R., Mekalanos J.J., Venter J.C.,
Fraser C.M.;
                                                    uncultured basidiomycete
Eukaryota; Fungi; Basidio
NCBI_TaxID=175244;
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TYGR; VC0279; -.
Hypothetical protein; Complete proteome.
SEQUENCE 42 AA; 4964 MW; 1C3B4EBDDB15BFEF CRC64;
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Submitted (NOV-1997) to the EMBL/GenBank/DDBJ databases
EMBL; Z99114; CAB13911.1; -.
Complete proteome.
                                                                                                                                                                                                            01-MAR-2002 (TrEMBLrel. 20, Creat
01-MAR-2002 (TrEMBLrel. 20, Last
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Laccase (EC 1.10.3.2) (Fragment).
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NCBI_TaxID=666;
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01-OCT-2000
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STRAIN-168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 390:249-256(1997).
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                                                                                                         Basidiomycota;
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Pred. No. 3e+02;
Mismatches
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   Luis P., Buscot F.;
"molecular biological monitoring of soil fungi with laccase genes.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ420344; CAD12471.1; -.
InterPro; IPR001117; Cu-oxidase.
Pfam; PF00394; Cu-oxidase; 1.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
Glucan 1,6-beta-glucosidase (EC 3.2.1.-) (EXO-1,6-beta-glucanase)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAINCOKE C11,
Jayus J., McDougall B.M., Seviour R.J.;
Jayus J., McDougall B.M., Seviour R.J.;
Puriffication and proprieties of a (1.6)-beta-glucanase from
Acremonium sp. strain OxF C11.",
Submitted (JAN-2000) to the SWISS-PROY data bank.
-!- CATALYTIC ACTIVITY: HYPROLYSIS OF SUCCESSIVE GLUCOSE RESIDUES
FROM 1,6-BETA-D-GLUCANS AND DERIVED OLICOSACCHARIDES.
-!- SUBCELLULAR LOCATION: EXTRACELLULAR.
                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE.

MEDLINE=96086028; PubMed=8534110;
Tamaru Y., Araki T., Amagoi H., Mori H., Morishita T.;
Tamaru Y., Araki T., Amagoi H., Mori H., Morishita T.;
"Purification and characterization of an extracellular beta-1,4-
mannanase from a marine bacterium, vibrio sp. strain MA-138.";
Appl. Environ. Microbiol. 61:4454-4458(1995).
SEQUENCE 20 AA; 2192 MW; DB14359E0F4C7FC4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Acremonium sp.
Eukaryota; Fungi; Ascomycota; mitosporic Ascomycota; Acremonium.
NCBI_TaxID=5045;
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                                                                                                                                                                                                                                                                                                                                        Bacteria; Proteobacteria; gamma subdivision; Vibrionaceae.
VCBI_TaxID=662;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20;
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                                                                                                                 47 AA; 5419 MW; 3F5F74F8A8802097 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence Update)
01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
Beta-mannanase (EC 3.2.1.78) (Fragment).
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Pred. No. 9.2e+02;
1; Mismatches 0;
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75.0%;
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Matches 4; Conserv
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Matches 3; Conserv
 TISSUE=MYCELIUM;
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Mus musoria. (Mouse).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Mammalia, Eutheria, Cetartiodactyla, Suina, Suidae, Sus.
NCBI_TaxID=9823;
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MEDLINE=96219641; PubMed=8639832;
MEDLINE=96219641; PubMed=8639832;
MEDLINE=96219641; PubMed=8639832;
Thibandeau K., Borche L., Soulillou J.P., Blanchard D.;
"Characterization of porcine platelet glycoproteins recognized ! human natural anti-gal' anti-bodies.";
Blood 87:4636-4642(1996).
SEQUENCE 20 AA; 2092 MW; C2A467776631A33E CRC64;
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                                                              Score 17; DB 3; Length 20;
Pred. No. 9.2e+02;
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01-WAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAY-2000 (TrEMBLrel. 13, Last annotation update)
Heterogenous nuclear ribonucleoprotein A2/B1 (Fragment).
HNRNP A2/B1.
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Roshon M.J., DeGregori J., Ruley H.E.;
"Gene trap mutagenesis of hnRNP A2/B1.";
Submitted (UNN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF073990; AAD29846.1; -.
                    F831A01126E04B61 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 940042C753673474 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UNA-2000 (TrEMBLrel. 14, Last annotation update)
135 Kpa glycoprotein/GP IIB homolog (Fragment)
Sus scrofa (Pig).
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Pred. No. 9.2e+02;
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20 AA; 2175 MW;
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Matches 3; Conservative
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14 GSAF 17
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RESULT 15
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Q9WIE1;
Q9WIE1;
01-NOV-1999 (TrEMBLrel. 12, C.
01-NOV-1999 (TrEMBLrel. 12, L.
01-DEC-2001 (TrEMBLrel. 19, L.
Defective RNA, strain T411.
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Q9EBRO;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Envelope glycoprotein (Fragment).
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MEDLINE=99190445; PubMed=10092024;

Ayllon M.A., Lopez C., Navas-Castillo J., Mawassi M., Dawson W.O.,

Guerri J., Flores R., Moreno P.;

"New defective RNAs from citrus tristeza virus: evidence for a

replicase-driven template switching mechanism in their generation.";

J. Gen. Virol. 80:817-821 (1999).

EMBL; Y18368; CAA77146.1; -.

SEQUENCE 24 AA; 2691 MW; 277F9A071D51235A CRC64;
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Nelson J.A.E., Baribaud F., Edwards T., Swanstrom R.;
"Patterns of Changes in Human Immunodeficiency Virus Type 1 V3
Sequence Populations Late in Infection.";
J. Virol. 74:8494-8501(2000).
EMBL; AF155905; AAG09947.1; -.
AIDS; Coat protein; Glycoprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1.
Viruees; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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Viruses; ssRNA positive-strand viruses, no DNA stage; Closteroviridae;
Closterovirus.
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13 GSAF 16
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on; der;	region peptide #3.				10740		Peptide #413 encod	Human peptide #387	Chicken ghrelin-li	Eel ghrelin-like G	Rainbow trout 20aa	Ghrelin-like growt	Human ghreiin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Human ghreiin þeþt	Human ghrelin pept	Human ghrelin pept	Human ghrelin pept	Synthetic Ghrelin		Human ghrelin pept	Human ghrelin pept	elin-like	Lin,		

growth hormone deficiency.

Rattus norvegicus.

Homo sapiens. Sus scrofa. Bos taurus.

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SHE CONTRACTION OF THE CONTRACTI 24-JUL-2000; 2000WO-JP04907

23-JUL-1999; 29-NOV-1999; 26-APR-2000; 20 ; 99JP-0210002. ; 99JP-0338841. ; 2000JP-0126623.

(KANG/) KANGAWA K.

Kangawa K, Kojima M, Hosoda Ξ Matsuo Ή, Minamitake

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of infant growth disorders -
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                                         New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis
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                                                                                                                                  Disclosure; Page 7; 210pp; Japanese.
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29-NOV-1999; 99JP-0338841.
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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are direlin homologues and are concarcatised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid has been substituted by a locompasses the unmodified peptides; the DNA encoding the peptides comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-like growth hormone secretagogue (GHS) core region sequence.
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Disclosure; Page 7; 210pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       growth hormone secretagogue; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SGIP; zsig33;
somatomedin-C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SGIP peptide of zsig33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB20100;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              anorectic; antidiabetic; therapy; somatotropin;
; nutritional absorption modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ξ
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                                                                                                                       /note=
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                                                /note=
                                                                                                                                                                                                                                                                                                                "optionally substituted by Ile, Val, Tyr in variant SGIP peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               "optionally substituted by Gly, Ala, Met in variant SGIP peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "optionally substituted by Gly, Ala, Met in variant SGIP peptides"
                      "optionally substituted by 
Tyr in variant SGIP peptid
                                                                                              "optionally SGIP peptid
                                                                                                                                                                                                                                              "optionally
Met, Pro in
                                                                                                                                                                                                                                                                                                                                                                                        "optionally substituted by Trp, Val, Ile in variant SGIP peption
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "optionally substituted by Ser, Ala, Met in variant SGIP peptides"
                                                                                                                                                                                              "optionally
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                           onally substituted peptides"
                                                                                                                                                                       Val in
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Pred. No.
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                                                                                                                                                                     substituted by Ala, variant SGIP peption
                                                                                                                                                                                                                                            substituted by Gly, Ala, variant SGIP peptides"
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0.097;
                 peptides"
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                                                                                                                  γģ
                                                                                                                                                             oy Ala, Gly,
peptides"
                                                                                                                                                                                                                                                                                                                                                                                     oy Trp, Tyr,
peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 10;
                                                                                                                  Дeр
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                                             гув,
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                                                                                                                                                                                                                                                                                                                                          Phe,
                                                                                                                                                                                                                                                                                                                                                                                                                    Leu,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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CXX VX
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KW ZSig
KW Ggest
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is that of novel peptide fragment SGIP of zsig33 (see AAB20101), a previously described secreted protein that is transcribed in the gastrointestinal system. SGIP is a contract that is transcribed in the gastrointestinal system. SGIP is a contract the form of growth hormone secretagogue receptor, and is therefore useful for modulating secretion of growth hormone and insuling like growth factor 1. SGIP and its variant peptides, comprising residues 1-9, 2-9, 3-9, 4-9, 2-10, 3-10, 4-10, 3-11 or 4-11 of SGIP in which at least 1 residue may be substituted, are used in claimed methods for stimulating contractability in duodenum or jejunum contissue, modulating pancreatic secretion of hormones and digestive enzymes, inducing growth hormone secretion, and modulating gastric emptying. Amino acid substitutions in SGIP may result in agonist or antagonist activity. The SGIP peptides are also useful for raising antibodies, identifying the SGIP receptor, screening agonists and antagonists, and identifying cells, tissues or cell lines which respond to a SGIP-stimulated pathway.
                                                                                                                                                                                                                                                                                                                                                            Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
           zsig33; signal transduction; hormone; enzyme; neural developmel gastric contractility; nutrient uptake; digestive; pancreatic; insulin-like growth factor-I; growth hormone; bone; gastrointel glucose; osteopathic; anorectic; vulnerary; immunomodulator; G-protein coupled receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel variants of SGIP peptides for modulating contractility in duodenum or jejunum tissue, pancreatic secretion of hormones and digestive enzymes, inducing growth hormone secretion or modulations.
                                                                                                            Human zsig33 peptide epitope
                                                                                                                                                                             AAB62656;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gastric emptying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-123010/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sheppard
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                                                                                                                                             23-JUL-2001
                                                                                                                                                                                                             AAB62656 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                             μ
                                                                                                                                                                                                                                                                                                                   1 GSSFLSPE
                                                                                                                                                                                                                                                                                           GSSFLSPE
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                                                                                                                                                                                                                                                                                                                                                          100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0345157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; 11
/note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "optionally substituted by Gln, Asn, Ser, Thr, His, Ala in variant SGIP peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "optionally substituted by Asn Ser, Thr, His, Ala, Glu, Asp, Lys, Arg in variant SGIP peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Deisher TA,
                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                            huzsig33-2.
                                                                                                                                                                                                                                                                                                                                                          Score 40; DB Pred. No. 0.10; Mismatches
                                                                            hormone; enzyme; neural development;
                                                                                                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                          0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bishop PD;
                                                                                                                                                                                                                                                                                                                                                                                            BB
                                                                                                                                                                                                                                                                                                                                                                                           22;
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                                                                                                                                                                                                                                                                                                                                                                                           Length 11;
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                                               gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or modulating
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WO200138355-A2.

31-MAY-2001

22-NOV-1999;

Sheppard PO,

Homo sapiens.

Synthetic.

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WPI; 2001-159704/16.
                                                                                                                                                                                                  (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27 AA;
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                                                                                                          23-JUL-1999;
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                         01-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of forming a reversible peptide-
creceptor complex that involves providing an immobilized receptor, and
contacting the receptor with a zsig33 peptide (Comprising residues 24-37
of AAB65649), where the receptor binds to the zsig33 peptide. The method
is useful for purifying cells, purifying a peptide, stimulating signal
creamsduction in a cell expressing a receptor. It is also useful for
contractility, nutrient uptake, secretion of digestive and
contractility, nutrient uptake, secretion of digestive and
pancreatic enzymes and hormones, secretion of insulin-like growth factor
cl. secretion of non-zsig33 proteins. It is useful for modulating growth
chormone secretion in a mammal having a disease associated with abnormal
levels of growth hormone, such as osteoporosis, bone repair, bone
cremodeling, low osteoblast levels, cartilage repair and remodeling,
cremodeling, low osteoblast levels, cartilage repair and remodeling,
crediting and ovulation induction, treating a mammal having
metabolic disorder requiring neurological feedback, such as satiety
requilation, glucose absorption and metabolism and neuropathy-associated
gastroincestianal disorders, and stimulating glucose-induced insular
contractions and stimulating business and stimulating glucose-induced insular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               release in a mammal. The present sequence represents a human zsig33 peptide epitope, used to raise zsig33 anti-peptide antibodies.
                                                                                                                                                                                                                                                                                                                                                                    Forming reversible peptide receptor complex for purifying cell and peptides, stimulating signal transduction and modulating hormone secretion, involves contacting a receptor with zsig33 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Growth hormone secretagogue, GHS; ghrelin, calcium concentration elevation; infant growth disorder;
                                                                                                                                                                                                                                                                                     Bishop PD;
                                                                                                                                                                                                                                                                                   Deisher TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rat des-Gln14-ghrelin, SEQ ID NO:10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; Page 111; 111pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB60514 standard; peptide; 27 AA.
                                                                                                                                                 22-NOV-2000; 2000WO-US32074.
                                                                                                                                                                                             99US-0166765
                                                                                                                                                                                                                                                                                   Jaspers SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             growth hormone deficiency.
                                                                                                                                                                                                                                       (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8; Conservative
                                                                                                                                                                                                                                                                                                                            WPI; 2001-355879/37.
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Best Local Similarity
Matches 8; Conserv
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Rattus norvegicus

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WO200107475-A1.

1 GSSFLSPE 8 ||||||||| 1 GSSFLSPE 8

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Sequence

AAB60514;

RESULT 6

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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides overtors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptides. The peptides of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                                                                                                                                                                                           New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis of infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ó,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 40; DB 22; Length 27; 100.0%; Pred. No. 0.28; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Growth hormone secretagogue, GHS, ghrelin, calcium concentration elevation; infant growth disorder; growth hormone deficiency.
                                                                                                                                                                                                                     Matsuo H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human des-Gln14-ghrelin, SEQ ID NO:11.
                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 185; 210pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB60515 standard; peptide; 27 AA.
                                                                                                                                                                                                                     Hosoda H,
24-JUL-2000; 2000WO-JP04907,
                                                     99JP-0210002
                                                                              29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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Best Local Similarity 100.
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29-NOV-1999;
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 calcium concentration elevation; infant growth disorder; growth hormone deficiency.
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                                                                                                                                                                 23-JUL-1999; 99JP-0210002.
29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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WPI; 2001-159704/16
                                                 Kangawa K,
                                                                                                                                                                                                                                                                                                                                                                                            WO200107475-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Porcine des-Gln14-ghrelin, SEQ ID NO:17.
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                                                                                                            (KANG/) KANGAWA K.
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8; Conserv
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                                                    Kojima M,
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                                                    Новода
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                                                 Matsuo H,
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0.28;
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diagnosis
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characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
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  Sequence
                                                                                                                                                                                                                                                                                                                                                                   concentration characterised
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 189; 210pp; Japanese
27 AA;
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Query Match
Best Local (
                               Matches
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            1 GSSFLSPE 8
GSSFLSPE 8
                              Similarity 100.
8; Conservative
                                      100.0%;
                               0; Mismatches
                                       Score 40; DB 2
Pred. No. 0.28;
                                               22,
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                                              Length 27;
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RESULT 9
AAB60522
WO200107475-A1
                                    Bos taurus
                                                                      calcium concentration elevation; growth hormone deficiency.
                                                                                     Growth hormone secretagogue; GHS; ghrelin; calcium concentration elevation; infant growth
                                                                                                                                            Bovine ghrelin,
                                                                                                                                                                                  24-APR-2001
                                                                                                                                                                                                                                                        AAB60522 standard; peptide;
                                                                                                                                                                                  (first entry)
                                                                                                                                              SEQ ID NO:22
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01-FEB-2001
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23-JUL-1999;
29-NOV-1999;
26-APR-2000; 2
                                                                               24-JUL-2000; 2000WO-JP04907
(KANG/)
KANGAWA K.
                           2000JP-0126623
                                     99JP-0210002
99JP-0338841
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Kangawa K, Kojima M,

Hosoda

Ħ,

Matsuo H,

Minamitake

2001-159704/16

New peptide compounds which induce growth hormone secretion elevate cell calcium concentrations, useful in treatment and of infant growth disorders -Claim 3; Pagd 193; 210pp; Japanese and and diagnosis

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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid has been substituted by a slow encompasses the unmodified peptides; the DNA encoding the peptides cetors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth to accompanying side effects. The present sequence represents a chrelin-type growth hormone secretagogue (GHS) of the invention.
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27 AA; Seguence

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Gaps
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100.0%; Score 40; DB 22; Length 27; 100.0%; Pred. No. 0.28;
                        0; Indels
                         Mismatches
                        , 0
                          8; Conservative
Query Match
Best Local Similarity
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1 GSSFLSPE 8 σ 1 GSSFLSPE

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RESULT 10

AAG64943 standard; peptide; 28 AA. AAG64943; AAG64943

1

Neurone denaturation prevention method related peptide #5.

(first entry)

19-OCT-2001

Neurone denaturation; neurone death; growth hormone liberation inhibitor; cerebral infarction; oedema; Alzheimer's disease; Parkinson's disease; Pick's disease; dementia; amyotrophic lateral sclerosis; cancer; diabetic neuropathy; neuroprotective; antiinflammatory; nootropic; cytostatic.

Unidentified

/note= "modified by O(C=O)(CH2)6CH3" Location/Qualifiers /label= OTHER Key Modified-site

WO200147558-A1

05-JUL-2001

28-DEC-2000; 2000WO-JP09431

99JP-0375513 28-DEC-1999;

(KAKE) KAKEN PHARM CO LTD

Ξ Neo Fujita K, Σ Amakawa Ohyama T, Murata T,

WPI; 2001-536280/59.

ŏ Agents for treating diseases associated with denaturation or neurons comprise growth hormone liberation inhibitor ${\tt -}$

The present invention provides agents for treating or preventing diseases associated with denaturation or death of neurons, which comprise a growth hormone liberation inhibitor. These can be used for treating or

ö preventing diseases associated with denaturation or death of neurons including those due to cerebral isohaemic disorders such as cerebral infarction or oedema. Other causes of denaturation or death of neurons included Alzheimer's disease, Pick's disease, AIDS related dementia, Parkinson's disease, anyotrophic lateral sclerosis, diabetic neuropathy and anticancer treatments. The present sequence is a peptide described in the exemplification of the invention. Gaps ö 100.0%; Score 40; DB 22; Length 28; 100.0%; Pred. No. 0.29; Growth hormone secretagogue, GHS, ghrelin, calcium concentration elevation, infant growth disorder, growth hormone deficiency. Minamitake ö Matsuo H, 0; Mismatches AAB60508 standard; peptide; 28 AA. новода н, 23-JUL-1999; 99JP-0210002. 29-NOV-1999; 99JP-0338841. 24-JUL-2000; 2000WO-JP04907. 26-APR-2000; 2000JP-0126623 (first entry) Local Similarity 100. SEQ ID NO:2. Kojima M, WPI; 2001-159704/16. Rattus norvegicus. (KANG/) KANGAWA K. ğ æ WO200107475-A1. 28 1 GSSFLSPE 1 GSSFLSPE Rat ghrelin, 24-APR-2001 01-FEB-2001 Kangawa K, AAB60508; Seguence Query Match Matches AAB60508 88888888888 셤

The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion contentration in cells. The peptides are directlin homologues and are concarration in cells. The peptides are directlin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptides. The peptides invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention. New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis Claim 2; Page 180; 210pp; Japanese. infant growth disorders

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Sequence

28

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RESULT 12

AAB60509

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                                                                                                                                                           The invention relates to a novel peptide compound or its salt which . CC induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concentration in that at least one amino acid has been substituted by caracterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid has been substituted by calso encompasses the unmodified peptides; the DNA encoding the peptides; certors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the convention, and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases cof the invention are useful for promoting infant growth due to growth Darmone deficiency. The compounds of the invention are safe with con accompanying side effects. The present sequence represents a cc ghrelin-type growth hormone secretagogue (GHS) of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide compounds which induce growth hormone secretion elevate cell calcium concentrations, useful in treatment and
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29-NOV-1999;
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les 8; Conser
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100.0%; Score 40; DI
100.0%; Pred. No. 0.3
tive 0; Mismatches
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Pred. No. 0.29;
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RESULT 13
AAB60518
ID AAB60
XX AAB60
XX AAB60
XX AAB60
XX Calcd
XX Calcd
XX Growt
XX W0200
DD 01-PE
XX W29-NC
PR 29-NC
RANG
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                                                                                                                                                                                                                                                                                               The invention relates to a novel peptide compound or its salt which CC induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are CC concentration in cells. The peptides are ghrelin homologues and are CC characterised in that at least one amino acid has been substituted by a laso encompasses the unmodified peptides; the DNA encoding the peptides; compound. The invention CC also encompasses the unmodified peptides; the DNA encoding the peptides; cetors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by CC chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases CC of the invention are useful for treating and/or diagnosing diseases CC caused by a deficiency in growth hormone expression or activity. In CC particular, they are useful for promoting infant growth due to growth CC no accompanying side effects. The present sequence represents a CC ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                       Query Match
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29-NOV-1999;
26-APR-2000;
                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis
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1 GSSFLSPE 8
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8; Conserv
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; 99JP-0338841.
; 2000JP-0126623.
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Pred. No. 0.29;
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                                                                                                                                                                                 DB 22;
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RESULT 14 AAB60530

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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion condentration in cells. The peptides are gireful homologues and are concarration in cells. The peptides are gireful homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide. The peptides of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promocing infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
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                                                                                                                                   Growth hormone secretagogue; GHS; ghrelin; calcium concentration elevation; infant growth disorder;
                                                                                                     Dog ghrelin-like GH secretagogue peptide, SEQ ID NO:31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matsuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 197; 210pp; Japanese.
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AAB60530 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of infant growth disorders -
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29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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                                                                  (first entry)
                                                                                                                                                                     growth hormone deficiency.
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                                                                                                                                                                                                                                                                                                                                                                                                                    (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 8; Conserv
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The present invention relates to a truncated ghrelin analogue or their salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone secretion. Ghrelin agonist companies, facilitating arowth hormone deficient state, increasing muscle mass and bone density, treating sexual dysfunction in males or females, facilitating a weight gain, maintenance of wight; maintenance of physical functioning, recovery of physical function, and/or appetite increase, or appetite increase is particularly useful for a patient comparing a disease or disorder, or under going a treatment, accompanied by eight loss such as anorexia, bulind, cancer cachexia, acquired immune deficiency syndrome (AIDS), wasting, cachexia and wasting in frail clader) and samples of treatments accompanied by weight loss include chemotherapy, radiation therapy, tapmocrary or permanent immobilisation and dialysis, and ghrelin antegoniem is utilised to facilitate weight complications of diabetes including retinopathy, and/or cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular close, appetite decrease, weight maintenance, treat obesity, diabetes and criain about a weight loss can be used for e.g. to reduce the likelihood of such diseases and for treating such diseases. Ghrelin analogue induces growth hormone release from primary-culture pituiery cells in a dose-dependent manner without stimulating the release of other pituitary hormones. Unlike longer length ghrelin, ghrelin analogue can be companied analogue can be synthesised aasily and has increased solubility in physiological buffers.
                                              acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone, muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological; cardiant; litholytic; hepatotropic.
                         ghrelin analogue, growth-hormone secretagogue, GHS receptor, AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulimia, cancer, obesity, diabetes mellitus, hypertension, osteoarbhritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Page 34; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                        25-MAY-2001; 2001WO-US17026.
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Search completed: January 10, 2003, 15:59:13 Job time: 31.2727 secs 1 GSSFLSPE 8 1 GSSFLSPE 8 ð

Human ghrelin peptide analogue, compound 6.

(first entry)

21-MAY-2002

AAE19032;

AAE19032 standard; peptide; 28 AA

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Indels

100.0%; Score 40; DB 23; Similarity 100.0%; Pred. No. 0.29; 8; Conservative 0; Mismatches 0;

Query Match Best Local Similarity

Matches

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Sequence

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The present sequence is human ghrelin peptide analogue

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second envelope gl
second envelope gl
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                                                                                                               phycobilisome 99K
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Oy 2 Db 17	Query Matc Best Local Matches	RESULT 2 PC2337 second envelope glycopr C;Species: hepatitis C C;Date: 21-Mar-1995 #se C;Accession: PC2337 R;Kato, N.; Nakazawa, T Biochem. Biophys. Res. Biochem. Biophys. Residues: PC2 A;Accession: PC2337 A;Molecule type: genomi A;Residues: 1-27 <kat> A;Cross references: DDB A;Experimental source: C;Keywords: glycoprotei</kat>	Best Local Matches Qy 1 GS Db 16 GG	H H C C D D	RESULT 1 S03275 photosystem C,Species: S; C,Date: 01-D; C,Accession: R,Murata, N.		6 C C C C C C C C C C C C C C C C C C C
SSFLSP: TSFLSP	Match ocal Simis S 5;	hepat Mar-1 Mar-1 Naka Naka Naka Naka Naka Naka Naka Naka Naka PC2 1-27 1	Sim 5; SFLS: : NFLD	Partial Partia	tem II 22K B: Spinacia 01-Dec-1989 ion: S03275		21 21 21 21 21 21 21 20 20 20 20 20 20 20 20 20 20 20 20 20
7 22	larity Conserva	SULT 2 (2337 (ilarity (Conservat:	yn. kes. 1, 70 tial amino aci number: S03269 S03275 Spe: protein 1-25 <mur> y: chlorophyll chloroplast; p</mur>	: 22K protein - nacia oleracea :-1989 #sequence :03275 Kajiura, H.; Fu		\$50.00000000000000000000000000000000000
	65. 83. tive	irus uence , Miz , Miz , Miz of hu 34; M	146	7 P. O.F	- 5e		30 30 30 30 30 30 30
		gp70, revis .utani, .utani, .u. 206, .u. 207, .u. 209, .u. 209,	ິ ປາ ຝ _ື ຈະ	sequence sequence /b-bind otosynth			000000000000000000
	Score 26; DB 2; Len Pred. No. 63; 1; Mismatches 0;	2 arrelope glycoprotein, gp70, hypervariable region 1 bes: hepatitis C virus 21-Mar-1995 #sequence_revision 26-May-1995 #text_cl sion: PC2337 N.; Nakazawa, T.; Mizutani, T.; Shimotohno, K. Biophys. Res. Commun. 206, 863-869, 1995 Susceptibility of human T-lymphotropic virus type ence number: PC2334; MUID:95134269; PMID:7832798 sion: PC3337 ule type: genomic RNA ules: 1-27 <katp 1-27="" <kat<="" <katp="" td="" ules:=""><td>N</td><td>ces of the protein ding protein hesis; photosystem Score 29; DB 2;</td><td>(fragment) h) on 01-Dec-1989 # Y.; Miyao, M.;</td><td>ALIGNMENTS</td><td>\$71301 B97413 B97413 \$412846 \$41210 B82657 B82657 B81239 G81239 G81239 G81239 G97539 S23950 \$23950 \$4650 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500</td></katp>	N	ces of the protein ding protein hesis; photosystem Score 29; DB 2;	(fragment) h) on 01-Dec-1989 # Y.; Miyao, M.;	ALIGNMENTS	\$71301 B97413 B97413 \$412846 \$41210 B82657 B82657 B81239 G81239 G81239 G81239 G97539 S23950 \$23950 \$4650 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500 \$46500
	ıgth 2 Indel	n 1 - t_chan t_chan t_chan	Indels	s of pea	rt I		הַלֵּנָת מִּלִנִלת לא בּנֵּה א סִינּה ו
	27; Ls 0; Gaps 0;	t - hepatitis C virus (isolate E hange 08-Oct-1999 I infected cell line MT-2 to P 4.1; PID:d1008349; PID:g882084 days postinoculation cells	LB 0; Gaps 0;	a and spinach photosystem II (change 21-Aug-1998 a, T., Watanabe, A., Shinozaki,		ICL5 protein - Par hypothetical prote serine proteinase hypothetical prote for pig myosin-light-chain actin-related prot Ig heavy chain v r second envelope gl peroxisome prolife allophycocyanin li Ig heavy chain v r

RESULT 3

||: ||| |GSTVLSP 13

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Page 27

Best Loc Matches

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Deca-1,3-glucanase - kidney bean
C.Species: Phaseolus vulgaris (kidney bean)
C.Species: Phaseolus vulgaris (kidney bean)
C.Species: Phaseolus vulgaris (kidney bean)
C.Species: Domar-1997 #sequence_revision 19-Mar-1997 #text_change 13-Sep-1998
C.Accssion: S2023
R.Daugrois, J.H.; Lafitte, C.; Barthe, J.P.; Faucher, C.; Touze, A.; Esquerre-Tugaye, M. Arch. Bloochen. Blopphys. 292, 468-474, 1992
A.Acc. Bloochen. Blopphys. 292, 468-474, 1992
A.Acc. Bloochen. Bloochen. S20329; MUID:92117656; PMID:1731612
A.Steference number: S20329
A.Steference review of the process of the pr
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Cyspecies: Sus scrofa domestica (domestic pig)
Cyspecies: O3-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
CyAccession: PN0607
R/Sillard, R.; Joenvall, H.; Mutt, V.
Biochem. Biophys. Res. Commun. 195, 746-750, 1993
A/Title: Characterization of porcine intestinal cytochrome c oxidase subunit VIIc, puril
                                                                                                                                                                                                             Ciperacia Incomplaints. To caenorhaboltis elegans
Ciperacia Incorpaboltis elegans
Cipate: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
Cipate: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
Cipate: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
Cipate: 15-Oct-1999 #sequence_revision: 126748
Submitted to the EMBL Data Library, January 1998
A;Reference number: 220258
A;Reference propriet Indianary; translated from GB/EMBL/DDBJ
A;Residues: 1-20 - WILL
A;Residues: 1-20 - WILL
A;Residues: 1-20 - WILL
A;Residues: EMBL:AL021482; PIDN:CAB54436.1; GSPDB:GN00021; CESP:Y39A1B.4
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Pred. No. 2.7e+02;
1; Mismatches 2; Indels
                                                                                                                                                                                     hypothetical protein Y39AlB.4 - Caenorhabditis elegans
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A,Residues: 1-47 <SIL>
A,Experimental source: intestine
C,Superfamily: cytochrome-c oxidase chain VIIc
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83.3%;
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Best Local Similarity 57.1%;
Matches 4; Conservative
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Best Local Similarity
Matches 5; Conserv
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A, Map position: 3
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26 GSEYLDP 32
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A,Molecule type: genomic RNA
A,Residues: 1-27 «KAT»
A;Cross-references: DDBJ:D43650; NID:g882085; PIDN:BAA07765.1; PID:d1008350; PID:g882086
A;Experimental source: isolate B-2, inoculum
C;Keywords: glycoprotein
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J. Protein Chem. 10, 415-425, 1991
A;Title: Clostridium botulinum types A, B, Cl, and E produce proteins with or without A;Reference number: A44644; MUID:92143938; PMID:1781887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurotoxin-associated protein type B Hn+ 17K chain - Clostridium botulinum (fragment) C;Species: Clostridium botulinum C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-Oct-1994 C;Accession: B44644
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C:Species: Ictalurus punctatus (channel catfish)
C:Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 16-Aug-1996
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C'Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 1C'Accession: A28835
C'Accession: A28835
R'Lobb, C.J.; Olson, M.O.J.
J. Immunol. 141, 1236-1245, 1988
A;Title: Immunoglobulin heavy H chain isotypes in a teleost fish. A;Reference number: A92822; MUID:88285736; PMID:2456346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 27;
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A;Status: preliminary
A;Status: preliminary
A;Residues: 1-33 <SOM>
A;Note: sequence extracted from NCBI backbone (NCBIP:83799)
A;Note: 16-Val was also found
C;Keywords: hemagglutinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
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Pred. No. 78;
1; Mismatches
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A;Residues: 1-30 <LOB>
C;Keywords: heterotetramer; immunoglobulin
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71.4%;
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17 TSFLSP 22
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Matches

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protein-serine/threonine kinase - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Sep-1996 #sequence_revision 06-Sep-1996 #text_change
C;Accession: 138221; S37423
R;Schultz, S.J.; Nigg, E.A.
Cell Growth Differ. 4, 821-830, 1993
Cell Growth Differ. 4, 821-830, 1993
A;Reference number: 138211; MUID:94100173; PMID:8274451
A;Accession: 138211
                                             protein QA100006 - Arabidopsis thaliana (fragment) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 30-Jun-1992 #sequence_revision 06-Jan-1995 C;Accession: PA0027 R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A Bubmitted to JIPID, July 1994
                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 1-12 <SCH>
C;Keywords: chloroplast; photosynthesis; photosystem II; thylakoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            photosystem II 3.7K protein - spinach (fragment) C;Species: Spinacia oleracea (spinach) C;Date: 01-Dec-1989 #sequence_revision 01-Dec-1989 C;Accession: S01122 R;Schroeder, W.P.; Henrysson, T.; Akerlund, H.E. FEBS Lett. 235, 289-292, 1988
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A; Reference number: PA0001
A; Accession: PA0027
                                  A; Description: Separation and
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PA0027
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A,Accession: S01122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-48 < RES>
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Best Local
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80.0%;
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                                  characterization of Arabidopsis
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Pred. No. 1.8e+02;
1; Mismatches 0;
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Pred. No. 5
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J. 2e+02;
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                                                                                                    #text_change
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                                 proteins
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C;Date: vz via 19414
C;Accession: I49414
R;Ko, M.S.; Wang, X.; Horton, J
Mamm. Genome 5, 349-355, 1994
A;Title: Genetic mapping of 40
                                                                                                                                                                                                                                                                 A; Experimental & C; Genetics:
A; Gene: CP0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene CTLA-1 protein - western wild mouse (fragment)
C;Species: Mus spretus (western wild mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 22-Jun-1999
RESULT 14
$04627
$194627
glutathione transferase (EC 2.5.1.18)
C;Species: Proteus mirabilis
                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-37 <REA>
A;Cross-references: GB:AE002224; GB:AE002161; NID:g7189574; PIDN:AAF38475.1; PID:g718957;
A;Cross-references: strain AR39, HL cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein CP0663 [imported] - Chlamydophila pneumoniae (strain AR39) C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 11-May-2000 C;Accession: A81552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Gene: CTLA-1
C;Superfamily: trypsin; trypsin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Residues: 1-15 <KAM>
A;Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       , C.; Dodson, R.; Gwinn, M.; Nelson, W. Nucleic Acids Res. 28, 1397-1406, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, I. C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-21 < RES>
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                                                                                                                                                                                                                                                                                                                                                                                                A;Reference number: A81500;
A;Accession: A81552
                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39
A;Reference number: A81500; MUID:20150255; PMID:10684935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Cross-references:
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                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary
                                                                                                                                                                                         Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL:U05708; NID:g497037; PIDN:AAB60471.1; PID:g497038
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66.7%;
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1; Mismatches
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Pred. No. 6
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2.3e+02;
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                       mirabilis
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C;Accession: S04627 A.; Piccolomini, R.; Allocati, N.; Caccuri, A.M.; Barra, D.; Feder FBS Lett. C.; Aceto, A.; Piccolomini, R.; Allocati, N.; Caccuri, A.M.; Barra, D.; Feder A;Title: N-terminal region of Proteus mirabilis glutathione transferase is not homologou A;Reference number: S04627; MUD: 89290034; PMID: 2661269 A;Reference type: protein A;Residues: 1-38 < DI: C;Complex: dimer C;Complex:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypotherical protein BH0662 [imported] - Bacillus halodurans (strain C-125)
C;Species: Bacillus halodurans
C;Species: Bacillus halodurans
C;Species: Bacillus halodurans
C;Accession: F83732
R;Takami, H; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hira
Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Recession: F83732
A;Recession: F83732
A;Recession: F83732
A;Recession: GB:AP001509; GB:BA000004; NID:g10173176; FIDN:BAB04381.1; GSPDB:GNOG C;Genetics:
A;Reperimental source: strain C-125
A;Gene: BH0662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A,Description: catalyzes conjugation of glutathione to a large variety of electrophilic bic compounds; involved in detoxification of organic hydroperoxides C;Superfamily: glutathione transferase C;Keywords: dimer; transferase
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C,Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 01-Feb-1999
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Search completed: January 10, 2003, 15:56:29 Job time : 14.7273 secs

16 SFVPPE 21

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GenCore version (c) 1993 - 2003

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Result
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Maximum Match 100%
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Maximum DB
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9
    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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    41476328 residues
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          CSPA KLEPN
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CSPA CITFR
CSPA SHIFO
IAPP BOVIN
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IAPP SHEED
IAPP SHEED
ATPO-SOLTU
LHB3 RHOAC
CSPA TEREN
R22A MOUSE
R711 BOVIN
BLP2 BOMON
GT SERMA
R22A MOUSE
TXJB HADVE
TXJB HADVE
TXJB HADVE
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TXJB HADVE
TXJB HADVE
DHF5 ERWCI
PSBN CYAPA
PSBN CYAPA
PSBN ANAVA
PSBN PSA
                                                                                                                                                                                                                                                                                                                                                              SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3754
Q48493 klebsiella
Q44078 aeromonas s
Q44078 aeromonas s
Q46051 citrobacter
Q46654 enterobacter
Q56178 salmonella
Q53816 shigella bo
Q54170 shigella fl
P81104 pinus pinas
Q48408 bacteriopha
P81383 ophiophagus
P81383 ophiophagus
P82507 bos taurus
Q28605 ovis aries
P80504 stroeptococc
Q56922 yersinia en
P35285 mus musculu
P82911 bos taurus
P22416 serratia ma
P82227 hadronyche
P82226 hadronyche
P82226 hadronyche
P82216 serratia ma
P82227 hadronyche
P82226 hadronyche
P82226 hadronyche
P82219 porphyra pu
P80510 triticum ae
P48108 cyanophora
P58560 anabaena va
P28642 pisum sativ
P14112 bacteriopha
                                                                                                                                                                                                                                                                                                                                 Description
                                                                                    B aeromonas h
T aeromonas g
1 citrobacter
4 enterobacte
8 salmonella
6 shigella fl
4 pinus pinas
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"Detection and speciation of bacteria through PCR using universal
"Detection and speciation of bacteria through PCR using universal
major cold-shock protein primer oligomers.";
-1 Ind. whicrobiol. Biotechnol. 19:286-293 (1997).
-1 SUBGNIT: HOMODIMER (BY SIMILARITY).
-1 SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
-1 INDUCTION: IN RESPONSE TO LOW TEMPERATURE (BY SIMILARITY).
-1 SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD) FAMILY.
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PROSITE; PS00352; COLD SHOCK; 1.
Transcription regulation; DNA-binding; Activator.
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15-UUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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Major cold-shock protein (Fragment)
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MEDLINE=98101990; PubMed=9439003;
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Pfam; PF00313; CSD; 1.
ProDom; PD000621; Cold_shock; 1.
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HSSP; P15277; 1MJC.
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                                                                 major cold-shock protein primer oligomers.";
J. Ind. Microbiol. Biotechnol. 19:286-293(1997).

-- SUBUNI: HOMODIMER (BY SIMILARITY).

-- SUBCELLULAR LOCATION: (BY STAPPARIC) (BY SIMILARITY).

-- INDUCTION: IN RESPONSE TO LOW TEMPERATURE (BY SIMILARITY).

-- SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD) FAMILY.
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STRAIN=NCTC 6071, and NCTC 6081 / C.ballerapensis;
MEDLINE=98101990; PubMed=9439003;
MEDLINE= Crowart G.S.A.B.;
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Major cold-shock protein (Fragment).
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Pfam, PP00313, CSD, 1.
ProDom, PD000621, Cold_shock; 1.
SMART; SM00357, CSP; 1.
PROSITE; PS00352; COLD_SHOCK; 1.
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Transcription regulation; DNA-binding;
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                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=NCTC 10006;
MEDLINE=98101990; PubMed=9439003;
Francis K.P., Stewart G.S.A.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOMAIN
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updat
                                                                                                                                           InterPro; IPR002059; Cold_shock. Pfam; PF00313; CSD; 1.
                                                                                                                                                                  EMBL; U60034; AAC80238.1; -. HSSP; P15277; 1MJC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enterobacter
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Bacteria; Proteobacteria; gamma subdivision; I
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                                                                                                         PROSITE; PS00352; COLD_SHOCK; 1.
                                                                                                                     SMART; SM00357; CSP;
                                                                                                                                                                                                                                                                                                                                                                      "Detection and speciation of bacteria through PCR using universal
                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=548;
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HSSP; P15277;
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3; Conserv
  Similarity
3; Conserv
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   Conservative
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; AA; 4983 MW;
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                                                                                                                              Cold_shock; 1.
             55.0%;
37.5%;
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37.5%;
score 22; DB; Pred. No. 3.5e.
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Pred. No. 3
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                                                 5B38560260B9E935 CRC64;
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                         DB 1;
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  .5e+02;
leg 2;
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                        Length 46;
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  Indels
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RESULT 7
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Best Local Similarity
Matches 3; Conserv
Bacteria;
Shigella.
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SEQUENCE
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J. Ind. Microbiol. Biotechnol. 19:286-293(1997).
-!- SUBUNIT: HOMODIMER (BY SIMILARITY)
-!- SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
-!- INDUCTION: IN RESPONSE TO LOW TEMPERATURE (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD) FAMILY.
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15-JUL-1998 (Rel. 36,
15-JUL-1998 (Rel. 36,
16-OCT-2001 (Rel. 40,
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                                                 Shigella boydii
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15-JUL-1998 (Rel. 36,
16-OCT-2001 (Rel. 40,
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                                                                                                                                                                                                                              CSPA
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Transcription regulation; DNA-binding;
NON_TER 1 1
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                                                                                                   Major cold-shock
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SMART; SM00357; CSP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=NCTC 5742;
MEDLINE=98101990; PubMed=9439003;
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Pfam; PF00313; CSD; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Detection and speciation of bacteria through PCR using universal
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                            Proteobacteria;
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                                                                                                . 36, Last sequence update) . 40, Last annotation updat protein (Fragment).
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37.5%;
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Last annotation updat
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Pred. No. 3.5e+02
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                               subdivision;
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                               Enterobacteriaceae;
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P81104;
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                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 major cold-shock protein primer oligomers.";
J. Ind. Microbiol. Biotechnol. 19:286-293(1997).
-!- SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
-!- INDUCTION: IN RESPONSE TO LOW TEMPERATURE (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD) FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                             STAIN=NCTC 9327;
MEDLINE=98101990; PubMed=9439003;
Francis K.P., Stewart G.-SA.B., B.,
"Detection and speciation of bacteria through PCR using universal major cold-shock protein primer oligomers.";
J. Ind. Microbiol. Biotechnol. 19:286-293(1997).
-i- SUBGNIT: HOMODIMER (BY SIMILARITY).
-i- SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
-i- SUBCINIT: HOMODIMER TO LOW TEMPERATURE (BY SIMILARITY).
-i- SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD) FAMILY.
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MEDLINE=98101990; PubMed=9439003;
Francis K.P., Stewart G.S.A.B.;
"Detection and speciation of bacteria through PCR using universal
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Pred. No. 3.5e+02;
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Transcription regulation; DNA-binding; Activator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last_annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Mismatches
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                                                                                                                                                                                                                                                         EMBL; U60036; AAC80240.1; -.
HSSP; P15277; 1MJC.
INTERFO; IPR002059; Cold_shock.
Pfam; PF00313; CSD; 1.
ProDom; PD000621; CSD; 1.
SMART; SM00357; CSP; 1.
                                                                                                                                                                                                                                                                                                                                  SHOCK;
                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 37.5%; 3; Conservative
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                                                                                                                                                                                                                                                                                                                     SMART; SM00357; CSP; 1.
PROSITE; PS00352; COLD
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Q54170;
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Matches
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use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-aib.ch/announce/or send an email to license@isb-sib.ch).
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15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1999 (Rel. 36, Last sequence update)
15-JUL-1999 (Rel. 38, Last annocration update)
16-JUL-1999 (Rel. 36, Last annocration update)

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-1- FUNCTION: PPIASE THAT BINDS CALMODULIN (BY SIMILARITY).

-1- CATALYTIC ACTIVITY: CIS-TRANS ISOMERIZATION OF PROLINE IMDIC PEPTIDE BONDS IN OLIGOPEPTIDES.

-1- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS PROTEIN IS: 5.3, ITS MI IS: 72 kDa.

-1- SIMILARITY: BELONGS TO THE FKBP-TYPE PPIASE FAMILY.

INTEPPO; IPRO01179; FKBP-PPIASE.

PROSITE; PS00444; FKBP-PPIASE.

PROSITE; PS00444; FKBP-PPIASE.

PROSITE; PS00445; FKBP-PPIASE.

PROSITE; PS00445; FKBP-PPIASE.

PROSITE; PS00446; FKBP-PPIASE.

PROSITE: PS04466; FKBP-PPIASE.

PROSITE: PS04
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Plomion C., Costa P., Bahrman N., Frigerio J.-M.,
Plomion C., Costa P., Bahrman N., Frigerio J.-M.,
"Genetic analysis of needle proteins in maritime pine. 1. Mapping
dominant and codominant protein markers assayed on diploid tissue,
a haploid-based genetic map.",
Silvae Genetica 46:161-165(1997).
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MEDLINE=99274088; PubMed=10344291;
Costa P., Pionneau C., Bauw G., Dubos C., Bahrman N., Kremer A.,
Frigerio J.-M., Plomion C.;
"Separation and characterization of needle and xylem maritime pine
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Pred. No. 3.5e+02;
3; Mismatches 2; Indels
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                                                                                                                                                          1675 MW; 2B53999722277F3F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR02059; Cold_shock.
Pfam; PF00313; CSD; 1.
ProDom; PP000621; Cold_shock; 1.
SMART; SMO357; CSP; 1.
PROSITE; PS00352; COLD_SHOCK; 1.
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Similarity 37.5%;
3; Conservative
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HSSP; P15277; IMJC.
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RESULT 10
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ID GP54_BPSP1
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15-DEC-1998 (Rel. 3
15-DEC-1998 (Rel. 3
                                                                                                                                                                                                                            15-DEC-1998 (Rel. 37, Created)
15-UL-1999 (Rel. 38, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
L-amino acid oxidase (EC 1.4.3.2) (LAO) (LAO) (Fragment).
Ophiophagus hannah (King cobra) (Naja hannah).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria, Squamata, Scleroglossa, Serpentes, Colubroidea;
Elapidae, Elapinae, Ophiophagus.
pit v
Arch.
[2]
                 MEDLINE=94361525; PubMed=8080286;
Ponnudural G., Chung M.C.M., Tan N.-H.;
"Purlfication and properties of the L-amino acid
pit viper (Calloselasma rhodostoma) venom.";
Arch. Biochem. Biophys. 313:373-378(1994).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=98327781; PubMed=9657951;
Stewart C.R., Gaslightwala I., Hinata K., Krolikowski K.A.,
Needleman D.S., Peng A.S.-Y., Peterman M.A., Tobias A., Wei P.;
"Genes and regulatory sites of the 'host-takeover module' in th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Viruses; dsDNA viruses, SPO1-like viruses.
                                                                                                                                              TISSUE=Venom;
                                                                                                                                                                     SEQUENCE.
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Р813<del>8</del>3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical
SEQUENCE 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     terminal redundancy of Bacillus subtilis bacteriophage SPO1.";
Virology 246:329-340(1998).
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2; Mismatches
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2; Mismatches 2;
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RESULT 13
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Ahn M.Y., Lee B.M., Kim Y.S.;
Ahn M.Y., Lee B.M., Kim Y.S.;
"Characterization and cytotoxicity of L-amino acid o renom of king cobra (Ophlophagus hannah).";
Int. J. Biochem. Cell Biol. 29:911-919(1997).
-I- FUNCTION: HAS CYTOTOXIC ACTIVITY (BY SIMILARITY)
-I- CATALYTIC ACTIVITY: An L-amino acid + H(2)O + O(4)
- NH(3) + H(2)O(2).
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P82537;
Eukaryota; Metazoa;
Mammalia; Eutheria;
Bovidae; Bovinae; Bo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kieselbach T., Bystedt M., Schroeder W.P.;
Submitted (MAY-2000) to the SWISS-PROT data bank.
-!- SUBCELLULAR LOCATION: Chloroplast; within the
Chloroplast; Thylakoid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Thylakoid lumenal 17 kDa protein (P17) (Fragment).
Spinacia oleracea (Spinach).
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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CONFLICT 1 1 H
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SEQÜENCE 19 AA; 2298 MW; D
                                                                        Bos taurus (Bovine).
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15-JUL-1998 (Rel. 36,
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-!- SUBUNIT: HOMODIMER (PROBABLE)
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TISSUE=Venom;
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SIMILARITY: BELONGS TO THE FLAVIN MONOAMINE OXIDASE
STRONG, TO MOUSE FIG-1.
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       Bos.
                          Chordata; Craniata; Vertebrata; Cetartiodactyla; Ruminantia; Pec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50.0%;
57.1%;
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                                                                                                                  Last sequence update)
Last annotation update)
tide (Amylin) (Fragment)
                                                                                                                                                                                           Created)
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Pred. No. 5.9e+02;
1; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EBD6462064CB67FF CRC64;
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3.6e+02;
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                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: SELECTIVELY INHIBITS INSULIN-STIMULATED GLUCOSE
UTILIZATION AND GLYCOGEN DEPOSITION IN MUSCLE, WHILE NOT AFFECTING
ADIPOCYTE GLUCOSE METABOLISM.
-!- SIMILARITY: BELONGS TO THE CALCITONIN FAMILY.
                                                                                                  Albrandt K., Sierzega M.E., Mull E., Brady E.M.G.;
Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
-!- PUNCTION: SELECTIVELY INHIBITS INSULIN-STIMULATED GLUCOSE
UTILIZATION AND GLYCOGEN DEPOSITION IN MUSCLE, WHILE NOT AFFECTING
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Sukaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinee; Ovis.
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Pred. No. 6.3e+02;
3; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                         - !- SIMILARITY: BELONGS TO THE CALCITONIN FAMILY.
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5-JUL-1998 (Rel. 36, Last annotation update)
slet amyloid polypeptide (Amylin) (Fragment)
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InterPro; IPR001693; Calcitonin-like.
Pfam; PF00214; Calc CGRP IAPP; 1.
SMART; SM00113; CALCITONIN; 1.
PROSITE; PS00258; CALCITONIN; PARTIAL.
Hormone; Amyloid.
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Pfam; PF00214; Calc CGRP IAPP; 1.
SMART; SM00113; CALCITONIN; 1.
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Best Local Similarity
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11 ANFLAP 16
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PEPTIDE
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Q28605;
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-!- SUBUNIT: F-TYPE ATPASES HAVE 2 COMPONENTS, CF(1) - THE CATALYTIC CORE - AND CF(0) - THE MEMBRANE PROTON CHANNEL. CF(1) HAS FIVE SUBUNITS: ALPHA(3), BETA(3), GAMMA(1), DELTA(1), BESILON(1). CF(0) HAS THREE MAIN SUBUNITS: A, B AND C.
-!- SUBCELLULAR LOCATION: Mitochondrial.
-!- SIMILARITY: BELONGS TO THE ATPASE DELTA CHAIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=TUBEY;
MEDLINE=97077345; PubMed=8919912;
Jansch L., Kruft V., Schmitz U.K., Braun H.P.;
Jansch L., Kruft V., Schmitz U.K., Braun H.P.;
Jansch L., Kruft V., Schmitz U.K., Braun H.P.;
Jansch L., Kruft V., Schmitz U.K.,
Ele procein complexes of plant mitochondria.";
Plant J. 9:357-368(1996).
-!- FUNCTION: THIS PROTEIN SEEMS TO BE PART OF THE STALK THAT LINKS
CF(0) TO CF(1). IT ETTHER THANSWITS CONFORMATIONAL CHANGES FROM CF(0) INTO CF(1) OR IS IMPLICATED IN PROTON CONDUCTION.
-!- CATALYTIC ACTIVITY: ATP + H(2)O + H(+)(In) = ADP + phosphate +
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01-FEB-1996 (Rel. 33, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
ATP synthase delta chain, mitochondrial (EC 3.6.3.14) (Oligomycin sensitivity conferral protein) (OSCP) (Fragment).
Solanum tuberosum (Potato).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae, euasterids I; Solanales; Solanaceae; Solanum.
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PROSITE; PS00389; ATPASE DELTA; PARTIAL.
Hydrolase; AIP synthesis; CF(1); Hydrogen ion transport; Mitochondrion.
                                                                             ISLET AMYLOID POLYPEPTIDE.
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                                                                                                                                  CB5609394EB44C05 CRC64;
                                                                                                                                                                             Score 20; DB 1; I
Pred. No. 6.3e+02;
3; Mismatches 0;
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Job time : 8.27273 secs
PS00258; CALCITONIN; PARTIAL Amyloid.
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                                                                                                                                3300 MW;
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les 3; Conservative
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11 ANFLAP 16
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32 SAFMXP 37
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                                                                        PEPTIDE
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GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.

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Minimum DB seq length: 0 Maximum DB seq length: 50
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

16	15	14	13	12	11	10	9	8	7	o	ហ	4	ω	ผ	1	Result No.
26	26	26	26	26	26	26	26	26	26	26	26	26	26	27	28	Score
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Q9ilf5 hepatitis c	Q9ilf4 hepatitis c	Q9ilf3 hepatitis c	Q9ilf2 hepatitis c	Q9ilf1 hepatitis c	Q9ilf0 hepatitis c		Q9ile8 hepatitis c	Q9ile7 hepatitis c		Q9qh84 hepatitis c	O37295 hepatitis c	Q81195 hepatitis c	Q81194 hepatitis c	Q9kig0 streptomyce	Q9zhw6 enterococcu	Description

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Query Match Best Local Similarity Matches 5; Conser	NON_TER SEQUENCE	Activa	SMART;	PRINTS; ProDom;	Pfam;	HSSP;	EMBL; AF047608;	-!- SU	bacteria.";	"Conse	Kim W.	STRAIN=EF1-1; MEDLINE=98440	SEQUENCE	NCBI_TaxID=1351;	Entero	Bacter	Enterococcus	Major cold	01-MAR-2002	01-MAY-1999	9WHZ6O	8 1
ch 1 Simi 5;	M	tor; [SMOO3	PDOC	PF0031	P32081	MILARI AF0476	BCELLU	1a.";	rvatio	S., K	EEF1-1	CE FRO	axib=1	coccac	ia, Fi	coccus		-2002	-1999	••	
Conser	50 50 AA;	NA-bin	SMART; SM00357; CSP; 1.	0621;	Pfam; PF00313; CSD; 1.	P32081; 1CSP.	17Y: BE	TLAR LO	2	n of t	unajak	0326;	FROM N.A.	.351;	eae; E	Bacteria; Firmicutes;	faecalis	shock p	(TrEMBLrel.	(TrEMBLrel.	PRELIM	
70.0%; larity 62.5%; Conservative	50 5340	Activator; DNA-binding; Transcription regulation.	PROSITE: PS00357; CSP; 1.	PRINTS; PR00050; COLDSHOCK. ProDom; PD000621; Cold_shock; 1.); 1.]	-i- SIMILARITY: BELONGS TO THE COLD-SHOCK DOMAIN (CSD)	-!- SUBCELLULAR LOCATION: CYTOPLASMIC (BY	bacteria.";	"Conservation of the major cold shock protein	H N.	STRAIN=EF1-1; MEDLINE=98440326; PubMed=9767713;			Enterococcaceae; Enterococcus.			shock protein (Fragment).			PRELIMINARY;	
μ.	MW; /	Transc	HOCK:	hock;	r.	٠	10 THE	CYTO	11955	or col	Ren J	=97677			occus.	cillus	trepto	(Frag	20, La			
۲ C	163EFE:	riptic	-	1.	}	÷	. COTO	DPLASM:	9091	ld show	, Duni	713;				/Clost	coccus	gment)	Last and	Created)	PRT;	
ore 28; DB pd. No. 92; Mismatches	A63EFE118510CF1E	n regu					SHOCK	(BY		k prot	N.W.					Bacillus/Clostridium group;	faeca		Last annotation update)		50 1	
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Length 50; Indels	CRC64;						N (CSD	SIMILARITY).		n lactic									ate))_		
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NCBI_TaxID=11103;
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STRAIN-NRRIJ5009,

Neu J.M., Wright G.D.;

"Characterization of stoPK-1, a novel protein kinase from the glycopeptide antibiotic producer Streptomyces toyocaensis NRRIJ5009.";

Submitted (PEB-2000) to the EMBL/GenBank/DDBJ databases.

EMBL, AF234273, AAF82534.1.

InterPro: IPR000719; Euk_pkinase.

PFam; PF00069; pkinase; 1.

ProDom; PD000001; Euk_pkinase; 1.

PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
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Kato N., Nakazawa T., Mizutani T., Shimotohno K.;
"Susceptibility of human T-lymphotropic virus type I infected cell
line MT-2 to hepatitis C virus infection.";
Biochem. Biophys. Res. Commun. 206:863-869(1995).
EMBL. D43649; BAA07764.1;
Pfam; PF01560; HCV NSI; I.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein;
Polyprotein; Transmembrane.
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein (Contains: envelope glycoprotein E2 (GP68) (GP70)
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Hepacivirus.
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Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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                                                                                                                                                                                                                                             OSKIGO;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Protein kinase Pk-3 (Fragment).
                                                                                                                                                                                                                              49 AA
                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptomyces toyocaensis.
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                     GSSFLSPE 8
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44 SYLSPE 49
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EMBL, APO18404, AAC036911; -
InterPro; IPR002531; HCV_MS1.
                                                                                                           Gaps
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Genome polyprotein (Contains: envelope glycoprotein E2 (GP68) (GP70)
(NS1)] (Fragment).
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"Susceptibility of human T-lymphotropic virus type I infected cell
line MT-2 to hepatitis C virus infection.";
Biochem. Biophys. Res. Commun. 206:863-869(1995).
EMBL; D43650; BAA07765.1; -.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein;
Polyprotein; Transmembrane.
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Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
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01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein (Contains: envelope glycoprotein E2 (GP68)
                                             Score 26; DB 12; Length 27; Pred. No. 1.3e+02; 1; Mismatches 0; Indels
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04170DDCFF3CAA7A CRC64;
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MEDLINE=95134269; PubMed=7832798;
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                                             65.0%;
2805 MW;
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Q9ILE6;
01-OCT-2000 (TrEMBLrel. 15, Cres
01-OCT-2001 (TrEMBLrel. 15, Lasi
01-DEC-2001 (TrEMBLrel. 19, Lasi
Genome polyprotein [Contains: ei
(NS1)] (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF01560; HCV_NS1; 1.
Coat protein; Envelope prote:
Polyprotein; Transmembrane.
NON_TER
NON_TER
27
SEQÜENCE
27 AA; 2757 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9QH84;
01-MAY-2000
01-MAY-2000
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein (Contains: envelope glycoprotein (NS1)) (Fragment).
                                                                                                                           Hepatitis C virus.
Viruses; ssRNA positive-strand
Hepacivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sandres K., Dubois M., Pasquier C., Izopet J.; "The genetic heterogeneity of the hypervariable genome and the sensitivity of hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus.
Hepatitis C virus.
Viruses; ssRNA positive-strand viruses, no DNA stage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polyprotein; Transmembrane.
NON_TER 1 1
NON_TER 27 27
SEQUENCE 27 AA; 2829 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR002531; HCV_NS1.
Pfam; PF01560; HCV_NS1; I.
Coat protein; Envelope protein; Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (JUL-1999) to the EMBL/GenBank/DDBJ EMBL; AF167044; AAD53677.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9QH84
                      Izopet J., R
Rumeau J.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                  SEQUENCE FROM N.A.
    Longitudinal analysis
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5; Conserv
                        Rostaing L
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83.3%;
, Sandres K., Cisterne J.M.,
., Durand D., Puel J.;
of Hepatitis C virus replica
                                                                                                                                                                                                                               Created)

Last sequence update)

Last annotation update)

s: envelope glycoprotein
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1; Mismatches
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Pred.
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                                                                                                                                                 viruses,
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  replication and liver
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                                          Pasquier
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RESULT

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ID Q9
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Best Local Similarity
Matches 5; Conserv
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Q9ILE7;
01-OCT-2000
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein (Contains: envelope glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fibrosis progression in renal transplant recipients."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases EMBL; AF207117; AAF75323.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Viruses; ssRNA positive-strand Hepacivirus. NCBI_TaxID=11103;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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NON_TER 1 1
NON_TER 27 27
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Pfam; PF01560; HCV_NS1; T.
Coat protein; Envelope protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis C virus
                                                      Hepatitis C virus.
Viruses; ssRNA positive-strand
                                                                                                                                                       (NS1)]
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(Fragment).
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2642 MW;
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83.3%;
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1; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3D471891B263D1CA CRC64;
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                                                             viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  viruses,
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                                                                Flaviviridae;
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Izopet J., Rostaing L., Sandres K., Cisterne J.M., Pasquier C., Rumeau J.L., Duffaut M., Durand D., Puel J.;
"Longitudinal analysis of Hepatitis C virus replication and liver fibrosis progression in renal transplant recipients.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.

EMBL, AR207114; AAF75320.1;
InterPro; IPR002531; HCV NS1.
Pram; PR01560; HCV NS1.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein;
Polyprotein; Transmembrane.
NON TER 1 1
NON TER 27 27
SEQÜENCE 27 AA; 2614 MW; 2FF0E891B273DOCA CRC64;
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PRELIMINARY; PRT; 27 AA.

Oglir]

Oll-OCT-2000 (TrEMBirel. 15, Created)

Ol-OCT-2000 (TrEMBirel. 15, Last sequence update)

Ol-OCT-2000 (TrEMBirel. 19, Last annotation update)

Ol-OFC-2000 (TrEMBirel. 19, Last annotation update)

Genome polyprotein [Contains: envelope glycoprotein E2 (GP68) (GP70)
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"Longitudinal analysis of Hepatitis C virus replication and liver fibrosis progression in remai transplant recipionts.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, AR207113; AAF7519.1;
- InterPro, IPR0025319.1;
- Fam: PR01560; HCV NS1.
Pfam: PR01560; HCV NS1.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein; PNON TER
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                 Hepatitis C virus.
Situses; BSRNA positive-strand viruses, no DNA stage; Flaviviridae;
Hepacivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis C virus.
Viruses, sBRNA positive-strand viruses, no DNA stage, Flaviviridae,
Hepacivirus.
NCBI_TaxID=11103;
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 26; DB 12; Length 27; Pred. No. 1.3e+02; 1; Mismatches 0; Indels
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Beet Local Similarity 83.33,
Beet Local Similarity 63.33,
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Best Local Similarity
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                                                                        NCBI_TaxID=11103;
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                     Izopet J., Rostaing L., Sandres K., Cisterne J.M., Pasquier C., Rumeau J.L., Duffaut M., Durand D., Puel J.;
"Longitudinal analysis of Hepatitis C virus replication and liver fibrosis progression in renal transplant recipients.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF207116; AAF75322.1;
InterPro: IPR002531; HCV NS1.
PFfam; PF01560; HCV NS1; I.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein; POlyprotein; Transmembrane.

NON TER 1 1
SEQÜENCE 27 AA; 2617 MW; 2FF0ED31B273D0CA CRC64;
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"Longitudinal analysis of Hepatitis C virus replication and liver fibrosis progression in renal transplant recipients.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, AF207115; ARF75221.1;
InterPro; IPR002531; HCV NS1.
Pfam; PF01560; HCV NS1.
Coat protein; Envelope protein; Glycoprotein; Nonstructural protein;
Polyprotein; Transmembrane.
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SEQÜENCE 27 AA; 2614 MW; 2FF0E891B273D0CA CRC64;
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O91LE9;
O1-CT-2000 (TrEMBLrel. 15, Last sequence update)
O1-OCT-2001 (TrEMBLrel. 15, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
O1-DEC-201 (TrEMBLrel. 19, Last annotation update)
(Senome polyprotein (Contains: envelope glycoprotein E2 (GP68) (GP70) (NS1)] (Fragment).
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2001 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein [Contains: envelope glycoprotein E2 (GP68) (GP70)
(NS1)] (Fragment).
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Hepacivirus.
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Pred. No. 1.3e+02;
1; Mismatches 0; Indels
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Q9ILF3,
Q9ILF3,
Q9ILF3,
C1-OCT-2000 (TrEMBLrel. 15, Created)
Q1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Q1-cma nolyprotein (Contains: envelope glycoprotein
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EMBL, AP207111; AAP75317.1; -. InterPro; IPR002531; HCV_NS1.

Pfam; PF01560; HCV_NS1; 1.
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"Longitudinal analysis of hepatitis C virus replication and liver fibrosis progression in renal transplant recipients.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, AF207112; AAF75318.1; -.
InterPro; IPR002531; HCV_NS1.
Pfam; PF01560; HCV_NS1; 1.
Coat protein; EnveTope protein; Glycoprotein; Nonstructural protein; Courterin; Transmembrane.
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Viruses, ssRNA positive-strand viruses,
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Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
Hepacivirus.
NCBI_TaxID=11103;
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Lrel. 19, Last annotation update)
[Contains: envelope glycoprotein E2 (GP68) (GP70)
                                                                                                                                                                                                                                                                                              2614 MW;
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Pred. No.
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                                                                                                                                                                                                                                                                                          2FF0E891B273D0CA CRC64;
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01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Genome polyprotein [Contains: envelope glycoprotein E2 (GP68) (GP70)
(NS1)] (Fragment).
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                                                                                                                                                                       Izopet J., Rostaing L., Sandres K., Cisterne J.M., Pasquier Rumeau J.L., Duffaut M., Durand D., Puel J.;
"Longitudinal analysis of Hepatitis C virus replication and fibrosis progression in renal transplant recipients.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, AF207110; AAF75316.1;
InterPro; IPR002531; HCV NS1.
Pfam; PF01560; HCV NS1; T.
Coat protein; Envelope protein; Glycoprotein; Nonstructural
                                                                                                                  SEQUENCE
                                                                                                                                                             Polyprotein; Transmembrane.
                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus
                                                                                                                                                                                                                                                                                                               EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                       iruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
17
                           2 SSFLSP 7
                                                                                                                                                                                                                                                                                                                                         _TaxID=11103;
ASFLSP 22
                                                       Similarity 5; Conserv
                                                                                                                  27 AA;
                                                          Conservative
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                                                                                                                 2642 MW;
                                                                     65.0%;
                                                      Score 26; DB 12;
Pred. No. 1.3e+02;
1; Mismatches (
                                                                                                                 3D471891B273D0CA CRC64;
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                                                                                   Length
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Minimum DB
Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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A Geneseq 101002:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES		

Regult No.	Score	Query Match	Query Match Length DB	80	ID	Description
ב	34	68.0	14	23	AAU76321	Synthetic Ghrelin
N	31.5	63.0	27	22	AAB60514	Rat des-Gln14-ghre
w	31.5	63.0	27	22	AAB60515	Human des-Gln14-gh
4	31	62.0	28	22	AAG64943	Neurone denaturati
v	1 2	62.0	28	22	AAB60508	Rat ghrelin, SEQ I
σ	31	62.0	28	22	AAB60509	Human ghrelin, SEQ
7	31	62.0	28	22	AAB60530	Dog ghrelin-like
8	31	62.0	28	23	AAE19032	Human ghrelin pept
9	31	62.0	28	23	AAE19040	
10	31	62.0	28	23	AAE19041	

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AAU76321; 21-MAY-2002 Synthetic Gh Ghrelin; ant acromegaly; Synthetic. Key Modified-sit WO200208250- 31-JAN-2002. 10-JUL-2001; 24-JUL-2000; (ZENT-) ZENT Deghenghi R; WPI; 2002-20	111 31 62 112 30 60 113 30 60 114 30 60 115 30 60 117 30 60 119 30 60 120 30 60 221 30 60 224 30 60 225 30 60 226 30 60 227 30 60 228 56 30 60 229 29 58 28 29 58 31 28.5 57 32 28 56 33 28 56 34 28 56 34 28 56 35 28 56 36 38 58 58 37 28 56 38 28 56 39 29 58 30 60 27 30 60 27 30 60 27 55 44 2 27 54 44 27 54 45 27 54 47 27 54 48 56 39 58 56 30 60 56 31 28 56 32 28 56 33 28 56 34 27 54 44 27 54 45 27 54 46 27 54
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entry) riant #3. growth h hic. tion/Qual el= OTHER e= "Other e= "Other 220178P.	peptide;
release; noyl Ser"	AAY40027 AAU99715 AABE9026 AAEE19021 AAEE19029 AAEE19030 AAEE19030 AAEE19033 AAEE19033 AAEE19037 AAEE19036 AAEE19037 AAEE19037 AAEE19038 AAEE19039 AAE
octanoyl ester; tumour;	Peptide sequence d Human Ghrelin (C-t Rat ghrelin-derive Rat ghrelin pept Human imelin pept Human imelin pept Human indigin ghaz-TH-IGG juncti Novel human diagno Biologically activ Bio

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invention.
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antagonises the effect of ghrelins when administered which to a mammal

Claim 3; Page 6; 9pp; English.

ghrelins when administered to a mammal. The peptide is a synthetic analogue of ghrelin. Ghrelin is a 27-28 residue peptide isolated from analogue of ghrelin. Ghrelin is a 27-28 residue peptide isolated from distinctive cell types in the stomach of rats and humans and has an octanoyl ester attached to a serihe residue. Ghrelin is a potent releaser of growth hormone. The peptides are useful for normalising elevated growth hormone levels in mammals such as those suffering from a tumour related to overproduction of growth hormone or acromegaly. The present sequence is a ghrelin antagonising peptide of the 늉 The invention relates to a novel peptide which antagonises the effect

14 AA; Sequence

Gaps 4; Score 34; DB 23; Length 14; Pred. No. 3.2;); Mismatches 1; Indels ö 68.0%; 64.3%; 9; Conservative 14 1 GSSF----AKLOPR 10 1 GSXFLSPEAKLOPR Query Match Best Local Similarity Matches ò g

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AAB60514 standard; peptide; 27 AA

24-APR-2001 (first entry)

Rat des-Gln14-ghrelin, SEQ ID NO:10.

Growth hormone secretagogue, GHS, ghrelin, calcium concentration elevation, infant growth disorder, growth hormone deficiency.

Rattus norvegicus

WO200107475-A1.

01-FEB-2001

24-JUL-2000; 2000WO-JP04907.

29-NOV-1999; 99JP-0338841. 26-APR-2000; 2000JP-0126623. 99JP-0210002 23-JUL-1999;

(KANG/) KANGAWA K.

Minamitake Y; Hosoda H, Matsuo H, Kangawa K, Kojima M,

WPI; 2001-159704/16.

New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis of infant growth disorders -

Claim 3; Page 185; 210pp; Japanese.

The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides, vectors and host cells comprising such DNA; a method of producing the

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                                       The peptides
               chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptid of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    elevate cell calcium concentrations, useful in treatment and diagnosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Growth hormone secretagogue, GHS, ghrelin; calcium concentration elevation; infant growth disorder; growth hormone deficiency.
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peptides comprising recombinant production, optionally
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                                                                                                                                                                                                                        63.0%; Score 31.5; Interity 37.0%; Pred. No. 21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human des-Gln14-ghrelin, SEQ ID NO:11.
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29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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les 10; Conserv
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                                                                                                                                                                                     27 AA;
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    Query Match 62. Best Local Similarity 35. Matches 10; Conservative
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Best Local
                                                                                                                                                    The present invention provides agents for treating or preventing diseases associated with denaturation or death of neurons, which comprise a growth hormone liberation inhibitor. These can be used for treating or preventing diseases associated with denaturation or death of neurons including those due to cerebral ischaemic disorders such as cerebral infarction or oedema. Other causes of denaturation or death of neurons included Alzheimer's disease, Pick's disease, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neurone denaturation; neurone death; growth hormone liberation innibitorerebral infarction; oedema; Alzheimer's disease; Parkinson's disease; Pick's disease; dementia; amyotrophic lateral sclerosis; cancer; disease; dementia; neuroprotective; antiinflammatory; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 17; 50pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                       Agents for treating diseases associated with denaturation or neurons comprise growth hormone liberation inhibitor -
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                                                                                                                                           invention.
  Score 31; DB Pred. No. 28; 0; Mismatches
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Pred. No. 21;
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                                               Length 28;
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Gaps
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                                                                                                                                                                                                                                                                        The invention relates to a novel peptide compound or its salt which CC induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are CC characterised in that at least one amino acid has been substituted by CC a modified amino acid and/or a non-amino acid has been substituted by CC also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by CC chemical modification; an antibody specific for a peptide of the invention are useful for treating and/or diagnosing diseases CC of the invention are useful for treating and/or diagnosing diseases of the invention are useful for promoting infant growth due to growth CC caused by a deficiency in growth hormone expression or activity. In CC particular, they are useful for promoting infant growth due to growth CC no accompanying side effects. The present sequence represents a CC ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                             Query Match
Best Local :
                                                                                                                     Matches
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29-NOV-1999;
26-APR-2000;
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                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; Page 180; 210pp; Japanese.
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                                                                                                                                          Local
                                                            1 GSSF-----
GSSFLSPEHQKAQQRKESKKPPAKLQPR
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                                                                                                                                             Similarity
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                                                                                                                  Conservative
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                                                                                                                                                                                                                                 ΑĄ,
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99JP-0338841.
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                                                      ----AKLOPR 10
                                                                                                                                          62.0%;
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                                                                                                               Score 31; DB
Pred. No. 28;
0; Mismatches
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28
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RESULT 6

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28 AA;
                                                        Canis familiaris
                                                                               WO200107475-A1
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                                                                                                                                                              29-NOV-1999;
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                                                                                                      01-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                New peptide compounds which induce growth hormone secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62.0%; Score 31; DB 22; Length 28; 35.7%; Pred. No. 28; ive 0; Mismatches 0; Indels
                                                                                       Growth hormone secretagogue, GHS; ghrelin; calcium concentration elevation; infant growth disorder; growth hormone deficiency.
                                                                                                                                                                                                                                                                                                   Minamitake
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dog ghrelin-like GH secretagogue peptide, SEQ ID NO:31.
                                                                                                                                                                                                                                                                                                   Matsuo H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -- AKLOPR 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GSSFLSPEHQRVQQRKESKKPPAKLQPR 28
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                                                                                                                                                                                                                                                                                                   Новода Н,
AAB60509 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                                     of infant growth disorders -
                                                                                                                                                                                                                               23-JUL-1999; 99JP-0210002.
29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
                                                                                                                                                                                                        24-JUL-2000; 2000WO-JP04907
                                                                  Human ghrelin, SEQ ID NO:3
                                            (first entry)
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                                                                                                                                                                                                                                                                                                   Kojima M,
                                                                                                                                                                                                                                                                                                                         WPI; 2001-159704/16.
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les 10; Conserv
                                                                                                                                                                                                                                                                            (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSSF-----
                                                                                                                                                            WO200107475-A1.
                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                              23-JUL-1999;
                                            24-APR-2001
                                                                                                                                                                                  01-FEB-2001
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                     AAB60509;
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Matches
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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concentration in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides ovectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptides. The peptides of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis
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Growth hormone secretagogue, GHS; ghrelin; calcium concentration elevation; infant growth disorder; growth hormone deficiency.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hosoda H, Matsuo H,
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 197; 210pp; Japanese.
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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone deficient state, increasing muscle mass and bone density, treating sexual dysfunction in males or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulimia, cancer, obesity, diabetes mellitus, hypertension,
21-MAY-2002
                                                                           AAE19040;
                                                                                                                                                  AAE19040 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a truncated ghrelin analogue or their salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 34; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAY-2000; 2000US-207920P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAY-2001; 2001WO-US17026.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cardiant; litholytic; hepatotropic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                       1 GSSFLSPEHORVQQRKESKKPPAKLQPR 28
                                                                                                                                                                                                                                                                                                                                                                                                                  1 GSSF----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
(first entry)
                                                                                                                                                  peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                  -----AKLOPR 10
                                                                                                                                                      28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31; DB
Pred. No. 28;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 23; Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1;
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BXAXAXB

Human ghrelin peptide analogue, compound 16.

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μ. Н

GSSFLXPEHORVOORKESKKPPAKLOPR

GSSF----

AKLOPR 10 0 Similarity

62.0%; 35.7%;

Score 31; Pred. No.

DB 28; 23; 0

Length Indels

18;

Gaps

Conservative

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CC females, facilitating a weight gain, maintenance of weight, maintenance of chysical functioning, recovery of physical function, and/or appetite increase is particularly useful for a patient companies, or appetite increase is particularly useful for a patient companies of the disorder, or under going a treatment, accompanied by eight loss such as anorexia, bulimia, cancer cachexia, acquired commune deficiency syndrome (AIDS), weathering, cachexia, and wasting in frail celderly and examples of treatments accompanied by weight loss include commotherapy, radiation therapy, temporary or permanent immobilisation and dialysis; and ghrelin antagonist is utilised to facilitate weight complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes weight is a contributing factor to different complications of such diseases including hypertension, diabetes, dyslipidemias, cardiovascular companies, osteoarthritis and certain forms of cancers, and contribution about a weight loss can be used for e.g. to reduce the companies in diseases including the contribution of the contribution of such diseases and for treating such diseases. Ghrelin collisions of such diseases and for treating such diseases of other pituitary hormones. Unlike longer length ghrelin, ghrelin analogue can be synthesised easily and has increased solubility in physiological buffers.

The present sequence is human ghrelin peptide analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone deficient state, increasing muscle mass and bone density, treating sexual dysfunction in males or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anoloulimia, cancer, obesity, diabetes mellitus, hypertension, osteoarthritis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ghrelin analogue; growth-hormone secretagogue; GHS receptor; AIDS; acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone; muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological; cardiant; litholytic; hepatotropic.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 23; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-195531/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-2000; 2000US-207920P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAY-2001; 2001WO-US17026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200192292-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to a truncated ghrelin analogue or their t, active at growth-hormone secretagogue (GHS) receptor. Ghrelin
28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Ser(CO-(CH2)6-CH3)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anorexia,
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1,

The present sequence is human ghrelin peptide analogue.

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acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis, growth hormone, muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological; cardiant; litholytic; hepatotropic.
                                                                                                                                                                                                                                                                                                                                                                             Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulimia, cancer, obesity, diabetes mellitus, hypertension,
                                                                             Human ghrelin peptide analogue, compound 17
                                                                                                                                                                                                                    /note= "Ser(CO-(CH2)6-CH3)"
                                                                                                                                                                                                 Location/Qualifiers
                  AAE19041 standard; peptide; 28 AA.
                                                                                                                                                                                                                                                                                                 30-MAY-2000; 2000US-207920P.
                                                                                                                                                                                                                                                                             25-MAY-2001; 2001WO-US17026.
                                                           (first entry)
                                                                                                ghrelin analogue;
                                                                                                                                                                                                                                                                                                                    (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-195531/25.
                                                                                                                                                                                                                                       WO200192292-A2
                                                                                                                                                                                                                                                                                                                                                                                                            osteoarthritis
                                                                                                                                                                                                         Modified-site
                                                                                                                                                                              Homo sapiens
                                                          21-MAY-2002
                                                                                                                                                                                                                                                           06-DEC-2001
                                                                                                                                                                                                                                                                                                                                        Bednarek M;
                                      AAE19041;
RESULT 10
          AAE19041
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growth-hormone secretagogue; GHS receptor; AIDS;

The present invention relates to a truncated ghrelin analogue or their salt, active at growth-hormons secretagogue (GHS) receptor. Ghrelin analogue is useful for screening a compound capable of binding to GHS caceptor and for stimulating growth hormons secretion. Ghrelin agonist is utilised for treating a growth hormone deficient state, increasing muscle mass and bone density, treating sexual dysfunction in males or females, facilitating a weight gain, maintenance of weight, maintenance of physical functioning, recovery of physical function, and/or appetite increase is particularly useful for a patient control of sorder, or under going a treatment, accompanied by increase, or appetite increase is particularly useful for a patient control of siscretar, or under going a treatment, accompanied by eight loss such as anorexia, bullmia, cancer cachexia, acquired by immune deficiency syndrome (AIDS), wasting, cachexia and wasting in frail contents accompanied by weight loss include conditions of treatments accompanied by weight loss include conditions of diabetes including retinopathy, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular consisting about a weight loss can be used for e.g. to reduce the bringing about a weight loss can be used for e.g. to reduce the lingining about a weight loss can be used for e.g. to reduce the lingining about a weight loss can be used for e.g. to reduce the lingining analogue induces growth hormone release from primary-culture pituitary cells in a dose-dependent manner without stimulating the release of other continutary hormones. Unlike longer length ghrelin, ghrelin analogue can be synthesised easily and has increased solubility in physiological buffers. Example 4; Page 23; 37pp; English.

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polynucleotides and their corresponding secreted polypeptides are useful for preventing, treating or ameliorating medical conditions, e.g. by protein or gene therapy. Pathological conditions can also be diagnosed by determining the amount of the new polypeptides in a sample or by determining the presence of mutations in the polynucleotide. Specific cancer, tumors, neurodegenerative disorders, development of and fetal deficiencies, blood disorders, scholognental abnormalities and fetal deficiencies, blood disorders, scholognents abnormalities system, autoimmune diseases, inflammation, allergies, Alzheimer's and cognitive disorders, entissue disorders, eschoporosis, arthritis, infections, AlDS, connective tissue disorders, transplant rejection, diabetes, asthma, sepsis, acne, psoriasis, cardiovascular disorders, and reproductive disorders. The polypeptides or polynucleotides can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human genes and the secreted polypeptides they encode, useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                                                                                                                                                                                                                                                                                            Secreted protein; gene therapy; cancer; tumor; fetal deficiency; neurodegenerative disorder; developmental abnormality; blood disorder; immune stylem disease; autoimmune disease; leukemia; inflammation; allergy; Alzheimer's disease; cognitive disorder; schizophrenia; obesity; osteoporosis; arthritis; infection; AIDS; diabetes; asthma; connective tissue disorder; respandar rejection; seppis; acceptatoria cardiovascular disorder; reproductive disorder; food additive; food preservative; storage capability.
                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rosen CA;
                                                                      18;
                                     Length 28
                                                                    0; Indels
                                                                                                                                                                                                                                                                                                             Peptide sequence derived from a human secreted protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ebner R, Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY40001-92 are derived from human secreted proteins.
                                    23;
                               DB ,
                                  Score 31; DB
Pred. No. 28;
0; Mismatches
                                                                                                   -----AKLQPR 10
                                                                                                                             1 GSSFLSPEHQRVQQRKEXKKPPAKLÖPR 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 27; 246pp; English.
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                                                                                                                                                                                                             AAY40027 standard; Peptide; 40
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98US-0076052.
98US-0076053.
98US-0076054.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US03939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0076057
                                                                                                                                                                                                                                                                              (first entry)
                                                                 Conservative
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                                                                                                  GSSF-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-550857/46.
                                                Similarity
 28 AA;
                                    Local Sim.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9943693-A1,
                                                                                                                                                                                                                                                                              18-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-SEP-1999.
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26-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-1998,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Olsen HS,
                                                                                                                                                                                                                                             AAY40027;
Sequence
                                Query Match
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                                                               Matches
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RESULT 12
AAU99715
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ភពពន្ធ
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                                                                                                                                                                                                                                                                                                                                01-NOV-2000;
29-MAY-2001;
19-OCT-2001;
The present invention describes a new method of treating a body weight disorder; increasing muscle mass and decreasing body fat by administration of angiotensin converting enzyme (ACE)-2 modulating compound. The invention can be used for treating body weight disorders, particularly obesity of at least grade 1, diabetes, atherosclerosis and a state associated with lipid metabolism. The method is used for treating rapid weight loss, rapid weight gain, anorexia, cachexia, bulimia, generalised partial lipodystrophy, familial partial lipodystrophy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; angiotensin converting enzyme-2; ACE-2; body weight disorder; muscle mass; body fat; obesity; diabetes; atherosclerosis; weight loss; lipid metabolism; weight gain; anorexia; cachexia; bulimia; sepsis; familial partial lipodystrophy; hypercholesterolaemia; hyperlipidaemia; aberrant metabolic rate; heart failure; left ventricular hypertrophy; neurodegenerative disorder; peptide hormone; cytokine processing; myocardial infarction; cardiomyopathy; inflammatory bowel disease; systemic inflammation response syndrome; polytrauma; pain; stroke; bone destruction; rheumatoid arthritis; osteoarthritis; asthma; periodontal disease; dymenorrhoea; premature labour; brain oddema; focal injury; diffuse axonal injury; reperfusion injury; scar formation; cerebral vasospasm; subarachnoid hearthries; altergic disorder; adult respiratory distress smallergic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     also be used as food additives or or decrease storage capabilities, carbohydrate, vitamins, minerals,
                                                                                                                                                                        Treating body administering
                                                                                                                                                                                                                       WPI; 2002-547572/58
                                                                                                                                                                                                                                                     Patane
                                                                                                                                                                                                                                                                                                                                                                                               31-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Ghrelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-SEP-2002
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                                                                                                                                           Example 18; Page 221; 395pp; English
                                                                                                                                                                                                                                                                                                    (MILL-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200239997-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                     e SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     respiratory distress syndrome; wound healing; appetite;
                                                                                                                                                                                                                                                                                                 MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40
                                                                                                                                                                                                                                                    Ocain TD, (
Kadambi VJ,
                                                                                                                                                                                                                                                                                                                              ; 2000US-0704216.
; 2001US-0870382.
; 2001US-371741P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US45703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                     weight disorder and increasing muscle mass comprises angiotensin converting enzyme-2 modulating compound
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ghrelin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               62.0%;
                                                                                                                                                                                                                                                                    Gould AE,
                                                                                                                                                                                                                                                     Solomon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide sequence.
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Pred. No.
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                                                                                                                                                                                                                                                                    Dales NA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preservatives, such as to increase fat content, lipid, protein, cofactors or other nutritional
                                                                                                                                                                                                                                                     Stricker-Krongrad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39;
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                                                                                                                                                                                                                                                                      Guan B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 40;
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                                                                                                                                                                                                                                                                    Brown
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AABGO SGO
ID AABGO
XX AABGO
AC AABGO
AC AABGO
AC AABGO
AC AABGO
XX 25-AP
XX Growt
KW Growt
KW Galci
KW Growt
XX Growt
KW Galci
XX CALCI
OS Synth
XX WO200
XX U0200
XX
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Matches
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                                                                                                                                                                                     New peptide compounds which induce g elevate cell calcium concentrations, of infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-1999;
29-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth hormone deficiency.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Growth hormone secretagogue; GHS; ghrelin; calcium concentration elevation; infant growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                myocardial infarction, cardiomyopathy, systemic inflammation response syndrome, sepsis, polytrauma, inflammatory bowel disease, acute and chronic pain, bone destruction in rheumatoid arthritis and osteoarthr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               congestive heart failure, chronic heart failure, left ventricular hypertrophy, acute heart failure, neurodegenerative disorders (e.g. Alzheimer's disease, harkinson's disease and Huntington's disease), diseases associated with peptide hormones or cytokine processing,
                                                                                                                                                                                                                                                                                                                                                       Kangawa K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUL-2000; 2000WO-JP04907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB60560;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB60560 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                   (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in the invention for hydrolysis of biologically active peptides
                                                                                                                                                                                                                                                                                                       2001-159704/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Conserv
                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 AA;
                                                                                                                                                                                                                                                                                                                                                       Kojima M,
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99JP-0338841
                                                                                                                                     84;
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                                                                                                                                     210pp;
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                                                                                                                                                                                                                                                                                                                                                             Hosoda
                                                                                                                                     Japanese
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                                                                                                                                                                                                                                              growth
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19;
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                                                                                                                                                                                                                          treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GHS) peptide
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The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted !

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ghrelin analogue; growth-hormone secretagogue; GHS receptor; AIDS; acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone; muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological; cardiant; litholytic; hepatotropic.
       also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides; opeptides comprising recombinant production, optionally followed by experience modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a truncated ghrelin analogue or their salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulimia, cancer, obesity, diabetes mellitus, hypertension,
                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                 ·;
                                                                                                                                                                                                                                                                            22; Length 28;
                                                                                                                                                                                                                                                                                                                 0; Indels
modified amino acid and/or a non-amino acid compound.
                                                                                                                                                                                                                                                                           60.0%; Score 30; DB 2
100.0%; Pred. No. 45;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Ser(CO-(CH2)6-CH3)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 3; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAE19021 standard; peptide; 28 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAY-2001; 2001WO-US17026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2000; 2000US-207920P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human ghrelin peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-195531/25
                                                                                                                                                                                                                                                                            Ouery Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                          28 AA;
                                                                                                                                                                                                                                                                                                                                                     5 AKLOPR 10
                                                                                                                                                                                                                                                                                                                                                                                       AKLOPR 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bednarek M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAE19021;
                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 14
   88888888888888888888
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is utilised for treating a growth hormone deficient state, increasing muscle mass and bone density, treating sexual dysfunction in males or females, facilitating a weight gain, maintenance of weight, maintenance of physical functioning, recovery of physical function, and/or appetite increase is particularly useful for a patient and adisease or disorder, or under going a treatment, accompanied by eight loss such as anorexia, bulina, cancer cachexia, acquired immune deficiency syndrome (AIDS), wasting, cachexia and wasting in frail eiderly and examples of treatments accompanied by weight loss include chemotherapy, radiation therapy, temporary or permanent immobilisation and dialysis; and ghrelin antagonist is utilised to facilitate weight complications of diabetes including retinopathy, and/or cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular chiseases, galls atones, osceoarthritis and certain forms of cancers, and bringing about a weight loss can be used for e.g. to reduce the consideration and one such an analogue induces growth hormone release from primary-culture pluniarry cells in a dose-dependent manner without strimlary abrelin analogue can be synthesised easily and has increased solubility in physiological buffers. ö acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone; muscle mass; bone density; sexual dysfunction; anorexia; wasting, radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological; cardiant; litholytic; hepatotropic. Gaps Human; ghrelin analogue; growth-hormone secretagogue; GHS receptor; ô 23; Length 28; 0; Indels /note= "Ser(CO-CH=CH-CH=CH-CH=CH-CH3)" The present sequence is human ghrelin peptide 60.0%; Scor. 100.0%; Pred. No. -c. Score 30; DB; Pred. No. 45; Human ghrelin peptide analogue, compound 1. Location/Qualifiers AAE19027 standard; peptide; 28 AA. 25-MAY-2001; 2001WO-US17026 30-MAY-2000; 2000US-207920P (first entry) Conservative (MERI) MERCK & CO INC Query Match Best Local Similarity 1 WPI; 2002-195531/25 28 AA; 28 5 AKLOPR 10 WO200192292-A2 Modified-site Homo sapiens 21-MAY-2002 06-DEC-2001. 23 AKLOPR Bednarek M; AAE19027; Sequence RESULT 15 Key AAE1902, 셤 ò

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The present invention relates to a truncated ghrelin analogue or their CC salt, active at growth hormone secretagogue (GHS) receptor. Ghrelin CC analogue is useful for screening a compound capable of binding to GHS receptor and for strimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone secretion. Ghrelin agonist CC muscle mass and bone density, treating sexual dysfunction in males or CC females, facilitating a weight gain, maintenance of weight, maintenance of freight loss such as anorexia, bulimia, cancer cachexia, acquired configurated as such as anorexia, bulimia, cancer cachexia, acquired configurated efficiency syndrome (AIDS) wasting, cachexia and wasting in frail celerly and examples of treatments accompanied by weight loss include chemotherapy, radiation therapy, temporary or permanent immobilisation complications of diabetes including retinopathy, and/or cachiovascular complications of diabetes including retinopathy, and/or cacdiovascular diseases including hypertension, diabetes, dysliphdemias, cardiovascular cdisease, gall stones, osteoarthritis and certain forms of cancers, and contrained by and has increased solubility in physiological buffers. The present sequence is human ahrelin peptide analogue can be synthesised easily and has increased solubility in physiological buffers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulimia, cancer, obesity, diabetes mellitus, hypertension,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; Page 33; 37pp; English
present sequence is human ghrelin peptide analogue.
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Sequence 28 AA;

ફ Matches Query Match Best Local (Local Similarity nes 6; Conserv 5 AKLQPR 10 Conservative 60.0%; Score 30; DB 23; Length 28; 100.0%; Pred. No. 45; tive 0; Mismatches 0; Indels <u>,</u> Gaps 0;

밁 Search completed: January 10, 2003, 15:59:14 Job time: 40.0909 secs 23 AKLQPR 28 THIS PAGE BLANK (USPTO)

GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 10, 2003, 15:55:16; Search time 15.9091 Seconds (without alignments) 60.427 Million cell updates/sec

Title: Perfect score: Sequence: C 50 1 gssfaklqpr 10

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

283224 seqs, 96134422 residues

Searched:

Total number of hits satisfying chosen parameters: 11827

Minimum DB seq length: 0 Maximum DB seq length: 50

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	ហ	4	w	N	_	
22	23	23	23	23	23	23	23	23	23	23	23	23	24	24	24	24	25	25	25	25	25	26	27	27	27	28	28	32	Score
44.0	6	46.0		ς,		46.0			σ.		რ	σ.	œ •		8	œ		0	0	0	0		4.	٠.	٠.	56.0	9		Query Match
13	50	45	44	43	40	39	37	36	30	29	27	19	50	48	39	30	49	43	29	25	11	39	37	27	27	19	16	43	Length
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PH1316	~	A81796	A60329	I45824	A99799	B81912	C70187	E41080	D81561	S02578	PQ0844	A34467	B82279	PQ0617	A47752	S08565	B97874	AI0982	B61613	S06263	86606	G89467	D83199	RHGPGA	RHPGA	T02624	I51879	S41388	
Ig heavy chain DJ		ical pr	PV9 -	C .	ical		het	_	χď	~	DNA-binding protei	36K microfibril-as	prot	DNA-directed DNA p	RNA recognition mo		degenerate transpo	hypothetical prote	ceratotoxin B - Me	gastrin-releasing	Q.	protein R09H3.2 (i	hypothetical prote	gastrin-releasing	gastrin-releasing	hypothetical prote	cystathionine beta	- hum	scription

RESULT 3

ALIGNMENTS

Qy 1 GSSFAKLQP 9 : : Db 8 GGAFAGLEP 16	Query Match 56.0%; Score 28; DB 2; Length 16; Best Local Similarity 55.6%; Pred. No. 14; Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps	RESULT 2 151879 Cystathionine beta-synthase - human (fragment) Cystathionine beta-synthase: 02-Jul-1996 #text_change 21-Jul-2000 Cystate: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jul-2000 Cystate: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jul-2000 Cystate: 02-Jul-1996 #sequence, M.P.; Panico, M.; de Franchis, R.; Kraus, J.P.; Al A, Title: The molecular basis of homocystinuria due to cystathionine beta-synthy. A; Status: preliminary; translated from GB/EMBL/DDBJ A; Status: preliminary; translated from GB/EMBL/DDBJ A; Cross-references: GB:S78267; NID:g999349; PIDN:AAB34404.1; PID:g999350	Query Match 64.0%; Score 32; DB 2; Length 43; Best Local Similarity 60.0%; Pred. No. 5.6; Matches 6; Conservative 2; Mismatches 2; Indels 0; Gaps Qy 1 GSSFAKLOPR 10 : : Db 29 GNPFAHLRPR 38	numan adenovirus 3 (fragment) stadenovirus h3 (human adenovirus 3) -1995 #sequence_revision 30-Jan-1998 # 541388 ; Chroboczek, J.; Jacrot, B. the EMBL Data Library, January 1994 : The penton base of human adenovirus tumber: \$41388 s41388 s41388 s41388 s41388 s41388 s41384 se: DNA -43 <cuz> ences: EMBL: Z29487; NID: g444048; PIDN: Cences: EMBL: Z29487; NID: g444048; PIDN: Cences: Serotype 3 is adenovirus peripentonal hexon-associa</cuz>	RESULT 1
	Gaps 0;	000 .; Andria -synthase	Gaps 0;	.ug-1999 9D motif. 1444049	
		defia			

deficie

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gastrin-releasing peptide - guinea pig
N/Contains: neuromedin C
C/Speciaes Cavia porcellus (guinea pig)
C/Speciaes Cavia porcellus (guinea pig)
C/Date: 31-Dec-1992 #sequence_revision 31-Dec-1992 #text_change 20-Mar-1998
C/Accession: A60206
C/Accession: A60206
C/Accession: A60206
A/Reference number: A60206; MUID:88034997; PMID:3668528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-27 <SHA>
A;Note: the carboxyl-terminal residue was not determined directly, but an alpha-amidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein PA3577 [imported] - Pseudomonas aeruginosa (strain PAO1)
C.Species: Pseudomonas aeruginosa
C.Species: Pseudomonas aeruginosa
C.Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000
C.Accession: D83199
R.Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; B. Adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim Nature 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Tille: Complete genome sequence of Pseudomonas aeruginosa PAO1, an opportunistic path. A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: D83139
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-37 <STO>
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C;Species: Caenorhabditis elegans
C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Nov-2001
C;Accession: G89467
R;anonymous, The C. elegans Sequencing Consortium.
R;anonymous, The C. elegans Sequencing Consortium.
A;Title: Genome sequence of the nematode C. elegans: a platform for investigating biolor A;Reference number: A75000; MUID:99069613; PMID:9851916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:AE004778; GB:AE004091; NID:g9949722; PIDN:AAG06965.1; GSPDB:GN00
A;Experimental source: strain PAO1
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S.Reywords: amidated carboxyl end; brain, hormone, intestine; neur
F.18-27/Product: neuromedin C #status predicted <NEU>
F:27/Modified site: amidated carboxyl end (Met) #status predicted
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Pred. No.
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62.5%;
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A,Status: translated from GB/EMBL/DDBJ
A,Status: translated from GB/EMBL/DDBJ
A,Molecule type: DNA
A,Resignes: 1-19 <ROU>
A,Cross-references: EMBL:AC004747; NID:g3413696; PID:g3413713
A,Experimental source: cultivar Columbia
A,Experimental source: cultivar Columbia
B,Ein, X.; Kaul, S.; Rounsley, S.D.; Blea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.;
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, L.
euss, D.; Nierman, W.C.; White, O.; Elsen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, J.
Aritle: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana.
A,Accession: B84655
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N/Contains: neuromedin C
C;Species us scrofa domestica (domestic pig)
C;Date: 31-Mar-1981 Hsequence revision 31-Mar-1981 #text_change 20-Mar-1998
C;Accession: A01562; A32738; C60301
C;Accession: A01562; A32738; C60301
C;Accession: A01562; A32738; C60301
Biochem. Biophys. Res. Commun. 90, 227-233, 1979
Biochem. Biophys. Res. Commun. 90, 227-233, 1979
A;Title: Characterization of a gastrin releasing peptide from porcine non-antral gastric A;Reference number: A01562; MUID:80042433; PMID:496973
                   hypothetical protein At2g25990 [imported] - Arabidopsis thaliana
NyAlternate names: hypothetical protein T19118.20
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 02-Feb-2001
C;Accession: T02624; B84655
C;Accession: T02624; B84655
Bshoulatey, S.D.; Kaul, S.; Lin, X.; Ketchum, K.A.; Crosby, M.L.; Brandon, R.C.; Sykes, submitted to the EMBL Data Library, August 1998
A;Description: Arabidopsis thaliana chromosome II BAC T19L18 genomic sequence.
A;Reference number: Z14681
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A; Residues: 1-27 < MCD>
A; Residues: 1-27 < MCD>
B; Minamino, N.; Kangawa, K.; Matsuo, H.
Biochem. Biophys. Res. Commun. 119, 14-20, 1984
A; Title: Neuromedin C: a bombesin-like peptide identified in porcine spinal cord.
A; Reference number: A32738; MUID:84153890; PMID:6546686
A; Accession: A32738
A; Molecule type: protein
A; Residues: 18-27 < MIN>
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A;Cross-references: GB:AE002093; NID:93413713; PIDN:AAC31236.1; GSPDB:GN00139
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1; Mismatches
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A,Gene: At2g25990; T19L18.20
A,Map position: 2
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quinoline 2-oxidoreductase alpha chain - Comamonas testosteroni (fragment) C;Species: Comamonas testosteroni (C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999 C;Accession: $66606 R;Schach, S.; Tshisuaka, B.; Fetzner, S.; Lingens, F. Bur. J. Biochem. 232, 536-544, 1995 A;Title: Quinoline 2-oxidoreductase and 2-oxo-1,2-dihydroquinoline 5,6-dioxygenase A;Reference number: $66606; MUID:96035889; PMID:7556204 A;Accession: $66606
                 RESULT
B61613
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A;Residues: 1-25 <CON>
A;Note: the sequence from the summary is
C;Superfamily: gastrin_releasing peptide
                                                                                                                                                                                                                                                                                                                         R;Conlon, J.M.; Henderson, I.W.; Thim, L. Gen. Comp. Endocrinol. 68, 415-420, 1987
A;Title: Gastrin-releasing peptide from the intestine of A;Reference number: S06263; MUID:88137922; PMID:3436516
                                                                                                                                                                                                                                                                                                                                                                                             gastrin-releasing peptide - smaller spotted catshark (tentative sequence) (fragment) C;Species: Scyllorhinus canicula (smaller spotted catshark, smaller spotted dogfish) C;Date: 31-Mar-1990 #sequence_revision 30-Jan-1998 #text_change 31-Mar-2000 C;Accession: S06263
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C; Keywords: neuropeptide
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C;Genetics:
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A; Residues: 1-39 < STO>
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A;Note: published errata appeared in Science 283, 35, 1999; Science 283, 2103, 1999; and
   ceratotoxin
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degenerate transposase (orf2) [imported] - Streptococcus pneumoniae (strain c,Species: Streptococcus pneumoniae
C;Species: Streptococcus pneumoniae
C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 22-Oct-2001
C;Accession: B97874
R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; Dei
e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; Mc
e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; Mc
y, P.; Sun, P.M.; Winkler, M.E.
J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jan
A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A;Reference number: A97872; MUID:21429245; PMID:11544234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein STY4163 [imported] - Salmonella enterica subsp. enterica (;Species: Salmonella enterica subsp. enterica serovar Typhi A;Note: this species has also been called Salmonella typhi (;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 09-Nov-2001 (;Accession: AI0882
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C;Date: 21-Unl-1995 #sequence_revision 28-Unl-1995 #text_change 17-Mar-1999
C;Accession: B61613
R,Marchini, D; Giordano, P.C.; Amons, R.; Bernini, L.F.; Dallai, R.
Insect Biochem. Mol. Biol. 23, 591-598, 1993
A;Title: Purification and primary structure of ceratotoxin A and B, two ant: A;Reference number: A61613; MUID:93357786; PMID:8353519
A;Accession: B61613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references:
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant
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C; Keywords: antibacterial;
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A;Molecule type: protein
A;Residues: 1-29 <MAR>
                                                                       A; Molecule type: DNA
A; Residues: 1-49 < KUR>
                                                                                                                                                 A;Reference number: A97872;
A;Accession: B97874
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A; Residues: 1-43 < PAR>
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                                             A;Cross-references: GB:AE007317; PIDN:AAK98822.1; PID:g15457547; GSPDB:GN00174
                                                                                                                       A; Status: preliminary
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1; Mismatches
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Pred. No. 1.9e+02;
3; Mismatches 1
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1.3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                D.; Wain, J.; Churcher,
L.; White, N.; Farrar,
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C;Accession: PQ0617

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NATIONAL TO SECONDITION MOTIF-type RNA-binding protein - fruit fly (Drosophila melanogaster) (from the companies of pages 24-Reb-1994 #sequence_revision 18-Nov-1994 #text_change 24-Sep-1999 (c) Accession: A47752 B.S. Mol. Cell. Biol. 13, 14-183, 1993 Mol. Cell. Biol. 14, RNA-type RNA-binding protein genes and the analysis of their relat A, Reference number: A48110; MuID: 93109300; PMID: 8417324 A, Accession: A47752 A, A, Status: preliminary A, Residues: 1-39 KINA A, Residues: 1-30 KINA A, Residues: 1-3
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R;Musgrove, J.E.; Johnson, R.A.; Ellis, R.J.
Rz. J. Biochem. 163, 529-534, 1987
A;Title: Dissociation of the ribulosebisphosphate-carboxylase large-subunit binding prot A;Reference number: S07232; MUID:87161853; PMID:3549295
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A;Cross-references: FlyBase:FBgn0010254
C;Superfamily: unassigned ribonucleoprotein repeat-containing proteins; ribonucleoprotei
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A,Residues: 1-30 <MUS>
C,Comment: This protein binds the newly synthesized large subunit and the as newly impos
C;Comment: This protein has ATPase activity.
C;Superfamily: chaperonin groEL
C;Keywords: chloroplast; heterododecamer; molecular chaperone
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DNA-directed DNA polymerase (EC 2.7.7.7) - human herpesvirus 6 (fragment)
C,Species: human herpesvirus 6
C,Date: 05-Aug-1994 #sequence_revision 05-Aug-1994 #text_change 07-May-1999
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                     Length 49;
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48.0%; Score 24; DB 2; Length 39;
Best Local Similarity 62.5%; Pred. No. 2.9e+02;
Matches 5; Conservative 1; Mismatches 2; Indels
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Score 25; DB 2; L
Pred. No. 2.2e+02;
1; Mismatches 1;
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71.4%;
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16 SAFAKYGP 23
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40 GSSFNKI 46
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R.Ellinger, K.; Naipel, F.; Foa-Tomasi, L.; Campadelli-Fiume, G.; Fleckenstein, J. Gen. Virol. 74, 495-500, 1993
J. M. Reference number: PQ0616; MUID:93187613; PMID:8383182
J. Reference number: PQ0616; MUID:93187613; PMID:8383182
J. Reference number: PQ0616; MUID:93187613; PMID:8383182
J. Residues: 1-48 ELL.
C, Superfamily: herpesvirus DNA-directed DNA polymerase
C; Keywords: DNA replication; nucleotidyltransferase
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seq length: 0 seq length: 50
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45.624 Million cell updates/sec
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    SwissProt_40:*
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             GRP PIG
PIG ADEB2
Q20A COWTE
GRP SCYCA
CERG CERCA
Y700 BORBU
ISP1 GALME
GRP ALLMI
Y00 BORBU
ISP1 GALME
GRP ALLMI
Y05 BPT7
LANG CARUI
SODM RANCA
RSHX THENCA
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TXAM MSTSE
OPZB-OXYKI
TRPD BACME
STYA STYCL
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P309472
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P31905
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P0321999
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P0112945
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P011295
P011477
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Q96627 bovine aden
P80464 comamonas t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description
                                                                                                                                5 metridium s
5 oxyopes kit
6 bacillus me
9 styela clav
0 styela clav
2 corynebacte
5 pleuronecte
6 spirulina p
7 thunnus obe
8 treponema p
6 haemophilus
                                                                                                                                                                                                                                                                                                4 commonas t
scyliorhinu
1 ceratitis c
3 borrelia bu
5 galleria me
6 alligator m
7 bacteriopha
0 carnobacter
5 rana catesb
3 thermus the
9 canis famil
7 amycolatops
3 fasciola he
5 gallus gall
0 rattus norv
1 borrelia bu
5 crotalus vi
9 crotalus vi
7 crotalus vi
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SQ

MOD_RES SEQUENCE

27 27 AA;

27 2806 MW;

AMIDATION. FA1017260210410F CRC64;

Query Match Best Local Similarity

54.0%;

Score 27; DB 1; Length 27; Pred. No. 23;

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3 L	nily; A	; Prozo44; Bom ITE; PS00257;	o; IPR000874;	RHGPGA	Dunca	Ξ,	ntestin	N: GRP	n. 49:1	ructure	im L.,	34997;	וולופוזיי	Julya.		C: a b	53890;	; FO-7;	10 . 31	орнув.	zation	Mutt V	:2:33; J., Joe			9823, 1	necazoa	llus (G	(Pia),				012	2			36.0	•	٠		•			
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ON C.						SIN/NEUROMEDIN B		release as well		ribution of guinea				14720(1304).		ide identified in				7-233 (1979) .	sing peptide from		G., Vaqne M.,				Suina: Suidae:				ence update) tation update)		27 AM.	1	MENTS	PL I	NA (100	EN	P1	, Ç	0;	3 2	
						/RANATENSIN		as othe		pig gast						porcine					porcin		Ghatei M.				Sus			n C (GRP-						P12801	P80652	P27751	Q56895	048408	P11860	P24330	P47910	
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SEQUENCE
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                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation. the Lucopean Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Quinoline 2-Oxidoreductase, alpha chain (EC 1.3.99.17) (Fragment).
Gomamonas testosteroni (Pseudomonas testosteroni).
Bacteria; Proteobacteria; beta subdivision; Comamonadaceae; Comamonas.
                                                                                                                                                                                                                     SEQUENCE FROM N.A.

Rusvai M., Harrach B., Banrevi A., Evans P., Benko M.;

Submitted (OCT-1996) to the EMBL/GenBank/DDBJ databases.

-i- FUNCTION: MINOR CAPSID PROTEIN THAT MAY ACT AS A LINK BETWEEN THE

EXTERNAL CAPSID AND THE INTERNAL DNA-PROTEIN CORE.
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-96035889; PubMed=7556204;
Schach S., Tshisuaka B., Fetzner S., Lingens F.;
"Quinoline 2-oxidoreductase and 2-oxor-1,2-dihydroquinoline 5,6-
quinoline and 3-methylquinoline degradation.";

Eur. J. Blacchen. 232:336-544(1995).
-!- FUNCTION: CONVERTS (3-METHYL-)-QUINOLINE TO (3-METHYL-)2-OXO-
1,2-DIHYDROQUINOLINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                    Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=114429;
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0
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 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                          BY SIMILARITY.
MINOR CAPSID PROTEIN VI.
                                                                                                                                                                                                                                                                                                                                                                                                                                          10F78E9678070306 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 26; DB 1;
Pred. No. 63;
2; Mismatches
                                                                                                                                                                           Bovine adenovirus type 2 (Mastadenovirus bos2)
                                                                                                                    Created)
Last sequence update)
Last annotation update)
                                                                                                                                                     Minor capsid protein VI precursor (Fragment).
                                                                                                  43 AA
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 Mismatches
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InterPro; IPR004243; McpVI.
Pfam; PF02993; MCPVI; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              52.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          4584 MW;
Conservative
                                                                                                STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                     (Rel. 35, C
(Rel. 35, I
(Rel. 35, I
                     1 GSSFAKLQPR 10
                                         8 GTVLAKMYPR 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | :|: | ||
3 GINFSALAPR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                          43 AA;
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01-NOV-1997
01-NOV-1997
 'n.
                                                                                                PIV6 ADEB2
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SEQUENCE
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Matches
Matches
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Gapa
-!- CATALYTIC ACTIVITY: Quinoline + acceptor + H(2)O = isoquinolin-
1(2H)-one + reduced acceptor.
-!- COPACTOR: FAD, MOLYBDENUM AND IRON-SULFUR.
-!- PATHMAX: FIRST STEP IN THE DEGRADATION OF QUINOLINE AND
(3-METHYL-)QUINOLINE.
-!- SUBUNIT: HETEROHEXAMER OF TWO ALPHA CHAINS, TWO BETA CHAINS, AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapa
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Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
dastrin-releasing peptide (GRP).
Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Blasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conlon J.M., Henderson I.W., Thim L.;
"Gastrin-releasing peptide from the intestine of the elasmobranch fish, Scyliorhinus canicula (common dogfish).";
Gen. Comp. Endocrinol. 68.415-420(1987).
-!- FUNCTION: GRP stimulates gastrin release as well as other gastrointestinal hormones.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: Secreted.
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Pred. No. 57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                             11 AA; 1213 MW; 869094322B1DC2CA CRC64;
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B735F911B89007F8 CRC64;
                                                                                                                                                                                           Oxidoreductase, Flavoprotein, FAD; Molybdenum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1994 (Rel. 29, Created) 01-JUN-1994 (Rel. 29, Last sequence update) 01-FB-1996 (Rel. 33, Last annotation update) Ceratoctoxin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; S06263; S06263.
InterPro; IPR000874; Bombesin.
Pfam; PF02044; Bombesin; 1.
PROSITE; PS00257; BOMBESIN; 1.
Bombeein family; Amidation.
MOD_RES 25 AMII
                                                                                                                                                                  TWO GAMMA CHAINS (PROBABLE)
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62.5%;
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Best Local Similarity
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Best Local :
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STRAINAATCC 35210 / B31;

MEDLINE=98065943; PubMed=9403685;

MEDLINE=98065943; PubMed=9403685;

Fraser C.M., Casjens S., Huang W.M., Sutton G.G., Clayton R.A.,

Lathigra R., White O., Ketchum K.A., Dodson R., Hickey E.K., Gwinn M.

Dougherty B., Tomb J.-F., Fleischmann R.D., Richardson D.,

Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M.,

van Vugt R., Palmer N., Adams M.D., Gocayne J.D., Weidman J.,

Utterback T., Watthey L., McDonald L., Artiach P., Bowman C.,

Garland S., Fujii C., Cotton M.D., Horst K., Roberts K., Hatch B.,

Smith H O. Venter T.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-DEC-1998
15-DEC-1998
16-OCT-2001
                                                                           This SWISS-PROT entry is copyright. It is produced through a collabora between the Swiss Institute of Bioinformatics and the EMBL outstati the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer entities requires a license agreement (See http://www.isb-sib.ch/annou or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibacterial peptides from the female reproductive accessory glands of the medfly Ceratitis capitata (Insecta Diptera)."; Insect Biochem. Mol. Biol. 23:1-598(1993).
-!- FUNCTION: FRANCE-SPECIFIC PROTIDES WITH POTENT ACTIVITY AGAINST GRAM-POSITIVE AND GRAM-NEGATIVE BACTERIA. THEY HAVE AS WELL HEMOLYTIC ACTIVITY. THESE PROTEINS ARE STABLE EVEN AT 100 DEGREES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical BB0700.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Female accessory gland;
MEDLINE=93357786; PubMed=8353519;
Hypothetical SEQUENCE 3
                                                                                                                                                                                                           Nature 390:580-586(1997).
                                                                                                                                                                                                                                                          Smith H.O., Venter J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Borrelia burgdorferi (Lyme disease spirochete).
Bacteria; Spirochaetales; Spirochaetaceae; Borrelia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BORBU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Purification and primary structure of ceratotoxin A and B, two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Marchini D., Giordano P.C., Amons R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE.
                               TIGR; BB0700;
                                              EMBL; AE001170; AAC67053.1;
                                                                                                                                                                                                                                      "Genomic sequence of a Lyme disease spirochaete, Borrelia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBUNIT: HOMOPOLYMER OF FOUR TO SIX SUBUNITS SUBCELLULAR LOCATION: Secreted.
SIMILARITY: STRUCTURALLY RELATED TO CECROPING
                                                                                                                                                                                                                                                                                                                                                                                                                                                  _TaxID=139;
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E 29 AA; 2861 MW;
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5; Conser
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pha; Tephritoidea; Tephritidae; Ceratitis.
l protein;
37 AA; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Rel. 37, Created)
(Rel. 37, Last sequence update)
(Rel. 40, Last annotation update)
protein BB0700.
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n; Complete proteome.
4267 MW; 39BAC907DE1B5B42 CRC64;
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55.6%;
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EE57F4EECB2DA6B0
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Pred. No.
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Best Local S
Matches 4
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MEDLINE=93324451; PubMed=8101369; Wang Y., Conlon J.M.; Wang Y., Conlon J.M.; "Neuroendocrine peptides (NPY, GRP, VII and stomach of the alligator."; Peptides 14:573-579(1993).
-!- FUNCTION: GRP stimulates gastrin regastrointescinal hormones.
-!- SUBCELLULAR LOCATION: Secreted.
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                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-1993 (Rel. 26, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Gastrin-releasing peptide (GRP) [Contains: Neuromedin C (GRP-10)].
Alligator mississippiensis (American alligator).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Crocodylidae; Alligatorinae; Alligator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P31886;
01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last seq
15-JUN-2002 (Rel. 41, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GRP_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Galleria mellonella (Wax moth).
Bukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda
Insecta; Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata;
Ditrysia; Pyraloidea; Pyralidae; Galleriinae; Galleria
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30-MAY-2000 (Rel. 39, Last sequence up
30-MAY-2000 (Rel. 39, Last annotation
Inducible serine protease inhibitor 1
                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=8496;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Serine protease inhibitor.
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                                                                                                                                                                                                                                                                                       SEQUENCE.
TISSUE=Stomach;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mellonella."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Hemolymph;
MEDLINE=20193629; PubMed=10727944;
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DEVELOPMENTAL STAGE: LAST INSTAR LARVAE.
INDUCTION: BY INPECTION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Biochem. 267:2046-2053 (2000)
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nilarity 44.4%;
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57.1%;
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No. 2.3e+02;
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STANDARD;

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SODM RANCA
P36215;
    CARUI
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                    P36960;
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                                                                                                                                                                                                                                                             Gaps
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Dunn J.J., Studier F.W.;
"Nucleotide sequence from the genetic left end of bacteriophage T7
"Nucleotide sequence from the genetic left end of bacteriophage T7
DNA to the beginning of gene 4.";
J. Mol. Biol. 148:303-330(1981).
-!- FUNCTION: THE FUNCTION OF THIS EARLY GENE PROTEIN IS UNKNOWN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteriophage T7.
Viruses; dsDNA viruses, no RNA stage; Caudovirales; Podoviridae;
T7-like viruses.
SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
                                                                                                                                                                                                                                                             1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=83241725; PubMed=6864790;
Dunn J.J., Studier F.W.;
Complete nucleotide sequence of bacteriophage T7 DNA and the locations of T7 genetic elements.";
J. Mol. Biol. 166:477-535(1983).
                                                                                                                                                                                                                45.0%; Score 22.5; DB 1; Length 28; 54.5%; Pred. No. 2.2e+02; Alsmatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44.0%; Score 22; DB 1; Length 47; 66.7%; Pred. No. 4.9e+02; tive 1; Mismatches 1; Indels
                                                                                                                                                   AMIDATION.
A74DB0487D844963 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
01-MAR-1989 (Rel. 10, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         47 AA
                                                                                                                               NEUROMEDIN C.
                                 InterPro; IPR000874; Bombesin.
Pfam; PF02044; Bombesin, 1.
PROSITE, PS00257; BOMBESIN; 1.
Bombesin family; Amidation. NE
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EMBL, V01127; CAA24329.1; --
PIR, A04402, W0BPS7.
PIR, S42285, S42285
SEQUENCE 47 AA, 4745 MW;
                                                                                                                                                   28 28
28 AA; 2786 MW;
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les 6; Conservative
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Best Local Similarity
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GASFGK 22
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SEQÜENCE
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P03777;
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VOS BPT7
VOS BPT8
VOS BACE
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RESULT 10 LANC_CARUI

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                                                                                                                                                                                                                                                                                                                                                     a Carnobacterium sp.";
Appl. Environ. Microbiol. 58:1417-1422(1992).
-i- FUNCTION: LANTHIONINE-CONTAINING PEPTIDE ANTIBIOTIC (LANTIBIOTIC),
ACTIVE ON GRAM-POSITIVE BACTERIA.
Antibiotic; Bacteriocin; Lantibiotic.
NON TER
                                                                                                                                                                                                                                                                   MEDLINE=92321768; PubMed=1622206;
Stoffels G., Nissen-Meyer J., Gudmundsdottir A., Sletten K., Holo H.,
                                                                                                                                                                                                                                                                                                                     Nes I.F.;
"Purification and characterization of a new bacteriocin isolated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gape
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01-JUN-1994 (Rel. 29, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Superoxide dismutase [Mn], mitochondrial (EC 1.15.1.1) (Fragment).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibla; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            h Similarity 50.0%; Score 21; DB 1; Length 7; Similarity 50.0%; Pred. No. 1.1e+05; 3; Conservative 3; Mismatches 0; Indels
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                                       01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Lantibiotic carnocin UI49 (Fragment).
7 AA.
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Pram; PF00081; aodfe; 1.
Probom; PD000475; SODismutase; 1.
PROSITE; PS00088; SOD_MN; PARTIAL.
NON dordoreductase; Manganese; Mitochondrion.
SEQUENCE 23 AA; 2594 MW; SDB0ED9B0E048
                                                                                                                                     Carnobacterium ap. (strain UI49)
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RESULT 13
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RSHX_THETH

P32143; P80383; O9F2A8;

P32143; P80383; O9F2A8;

T 01-OCT-1993 (Rel. 27, Created)

T 01-OCT-1993 (Rel. 27, Last sequence update)

T 15-JUN-2002 (Rel. 41, Last annotation update)

T 15-JUN-2002 (Rel. 41, Last annotation update)
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Best Local S
Matches 3
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MEDLINE-95045586; PubMed-7957245;
Tsiboli P., Herfurth E., Choli T.;
Tsiboli P., Herfurth E., Choli T.;
"Purification and characterization of the bacterium Thermus thermophilus.";
Eur. J. Biochem. 226:169-177(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Choli T., Franceschi F., Yonath A., Wittmann-Liebold B.; "Isolation and characterization of a new ribosomal protein from thermophilic eubacteria, Thermus thermophilus, T. aquaticus and
GRP_CANFA STANDARD;
P08989;
01-NOV-1988 (Rel. 09, Created)
                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                    Biol
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPECIES-T thermophilus; STRAIN-HB8 / ATCC 27634;
MEDLINE-21421773; PubMed-11530930;
Leontiadou F., Triantafillidou D., Choli-Papadopoulou T.;
"On the characterization of the putative S20-thx operon of Thermus
                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                       Local Similarity
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SIMILARITY: BELONGS TO THE S31E FAMILY OF RIBOSOMAL PROTEINS.
                                                                                                                                                                                                                                                          ; AJ295159; CAC15068.1; -. S33869; S33869.
                                                                                                   TYGKYRPR 23
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                                                                                                                                                           Conservative
                                                                                                                                                                                                                3206 MW;
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                                                                                                                                                        Score 21; DB 1;
Pred. No. 4.2e+02;
3; Mismatches 2
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01-OCT-1989
01-OCT-1989
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MOD RES
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1988 (Rel.
15-JUN-2002 (Rel.
                                                                                                                                           TISSUE=Cnidoblast;
Krebs H.C., Habermehl G.G.;
"Isolation and structural determination of
from the sea anemone Metridium senile.";
Naturwissenschaften 74:395-396(1987).
                                                                                                                                                                                                                           Metridium senile (Brown sea anemone) (Frilled sea anemone).
Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Metridiidae; Metridium.
                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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InterPro; IPR003582; ShKT.
SMART; SM00254; ShKT; 1.
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SIMILARITY: BELONGS TO
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                                                                                                              INHIBITORY TOXIN FAMILY
  SFAKLQ 8
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                                                                       Cnidocyst.
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(Rel. 12, Last sequence up
(Rel. 41, Last annotation
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. 41, Last
g peptide
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alsh J.H., Chew
                                                             3974 MW;
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                               42.0%;
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rtide (GRP) [Contains: Neuromedin C (GRP-10)].
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                      Score 21; DB
Pred. No. 6e+0
1; Mismatches
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9D9317261B7C7D65 CRC64;
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Pred. No. 4.4e+02;
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                                                              5ED9CC73509E007F
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ANEMONE POTASSIUM CHANNEL
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                                         Length 36;
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                       Indels
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Canis.
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RESULT 15

OPZB OXYKI STANDARD, PRT, 37 AA.

DD 0PZB OXYKI

DD 0PZB OXYKI

DD 0PZB OXYKI

DT 15-JUN-2002 (Rel. 41, Last sequence update)

DT 15-JUN-2002 (Rel. 41, Last sequence update)

DT 15-JUN-2002 (Rel. 41, Last sequence update)

OX OXPOPES Kitabensis (Wolf spider)

DT 15-JUN-2002 (Rel. 41, Last sequence update)

OC ENARYCER, Metazoa Arthropoda; Chelicerata; Arachnida; Araneae;

OX NCBI TAXIDIA 184771;

RN (1) TAXIDIA 184771;

RN (1) TAXIDIA 184771;

RN PECTROMETRY, AND CIRCULAR DICHROISM ANALYSIS.

RN PUBMEd-1197632;

RX PUBMEd-1197632;

RX PUBMEd-1197632;

RX PUBMED CANODER Kitabensis with cytolytic properties and positive for volginins, large amphipathic peptides isolated from the venom of the Novyopinins, large amphipathic peptides isolated from the venom of the ST 1976204 CANODER CANODER ST 197632;

RX PUBMED-1197632;

RX PUBMED-119763;

RX PUBMED-1197
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Search completed: January 10, 2003, 15:55:49 Job time : 11.0909 secs

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Minimum DB
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                                                                                                                                                                                                                                                                            Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seq length: 0
seq length: 50
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Gapop 10.0 , Gapext 0.5
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     SPTREMBL_21:*
1: sp_archea:
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sp_mammal:*
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sp_bacteria:*
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     GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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 P77066
P77498
P77514
P77055
Q21886
Q9R137
O89348
O88440
Q9IX54
Q8Z2A2
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Q16350
O80997
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P77063
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Q65289 human adeno
Q16350 homo sapien
O80997 arabidopsis
Q9hy43 pseudomonas
P7063 escherichia
P7064 escherichia
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P77067 escherichia
P77514 escherichia
P77514 escherichia
P77618 escherichia
P77619 escherichia
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Q8w4x5 arabidopsis	Q8W4X5	10	15		22	45
OS1559 borrelia bu	051559	16	50	•	23	44
irounga l	Q28491	6	50	σ,	23	43
_	Q991B9	12	46	46.0	23	42
Q9jsm4 neisseria m	Q9JSM4	16	45	ς,	23	41
Q44544 anabaena va	Q44544	N	44		23	40
Q28456 leptonychot	Q28456	σ	43	٥.	23	39
Q27942 arctocephal	Q27942	σ	43	9	23	38
Q28458 mirounga an	Q28458	σ	43	٥,	23	
ydrui	Q28444	σ	43	5	23	
hoca	Q28816	6	43	σ.	23	S
omo s	Q9H480	4.	43	٥,	23	34
esch	Q8X2K7	16	40	5	23	S S
neisseria	86nr6Ö	16	39	46.0	23	32
Ch1	Q9K239	16	30		23	31
mus mu	Q9CSC1	11	27	٥,	23	30
gal	Q8UVE1	13	26	ა.	23	29
equus	Q95MB1	თ	18	٥.	23	28
Q8uuh0 gallus gall	QBUUH0	13	13	46.0	23	27
vibr:	Q9KTT0	16	50	8	24	26
mue	Q8VIN5	11	48	æ	24	25
mus	Q8VBU3	11	41	æ	24	24
Bum	Q8VBX8	11	39	8	24	23
droso	Q26280	u	39	8	24	22
ana	Q8VBU6	11	35	8	24	21
	Q8VIN3	11	26	œ	24	20
hepatiti	056530	12	27	٥	24.5	19
rx6 xenopus	Q9PRX6	13	45		25	18
Q96cf1 homo sapien	Q96CF1	4	44	50.0		17

ALIGNMENTS

RESULT Q16350 ID Q AC Q DT 0	Qy Mat	S T T R R R R R R R R R R R R R R R R R	521
JT 2 30 Q16350 Q16350; Q1-NOV-1996 01-NOV-1996	Query Match Best Local Similar Matches 6; Con 1 GSSFAKLOPR : : 29 GNPFAHLRPR	[1] STRQUENCE FROM N STRAIN=SEROTYPE MEDLINE=9435744 Cuzange A., Chr "The pencon bass Gene 146:257-25 EMBL; Z29487; C InterPro; IPRO0 Pfam; PFO2455; NON TER SEQUENCE 43 A	1 65289; 1-NOV-1996 1-NOV-1996 1-JUN-2002 rotein 3a rotein 3a calenc
PRELIMINARY; (Tremblrel. 01,	Similarity 60.0%; Similarity 60.0%; 6; Conservative FFAKLOPR 10	.A. 3; 3; 3; 6; PubMed=8 6; PubMed=8 0boczek J., e oboczek J., p4194). p4194). p3479; Hex_I 14ex_IIIa; 1 14ex_IIIa; 1 14754 MW	PRELIMINARY; (TrEMBLrel. 01, (TrEMBLrel. 21, (TREMBLrel. 21, (TREMBLrel. 21, Pragment). Pragment). Irus type 3. Nirus type 3. Nirus type 3. 15659;
PRT;. 16 AA. Created) Last sequence update)	Score 32; DB Pred. No. 18; 2; Mismatches	076828; Jacrot B.; Jacrot B.; adenovirus type 3 h . IIa F8E42D960369585A	PRT; 43 AA. Created) Last sequence update) Last annotation update) RNA stage; Adenoviridae;
AA. . update)	12; Length 43;	as the	
	0; Gaps	RGD motif.";	Mastadenovirus.
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1 GSSFAKLO 8
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STRAIN=CV. COLUMBIA;
MEDLINE=20003487; PubMed=10617197;
Lin X., Kaul S., Rounsley S.D., Shea T.P., Benito M.-I., Town C.D.,
Lin X., Kaul S., Rounsley S.D., Shea T.P., Benito M.-I., Town C.D.,
Fujii C.Y., Mason T.M., Bowman C.L., Barnstead M.E., Feldblyum T.V.,
Buell C.R., Ketchum K.A., Lee J.J., Ronning C.M., Koo H., Moffat K.S.
Cronin L.A., Shen M., VanAken S.E., Umayam L., Tallon L.J., Gill J.E.,
Adams M.D., Carrera A.J., Creasy T.H., Goodman H.M., Somerville C.R.,
Copenhaver G.P., Preuss D., Nierman W.C., White O., Eisen J.A.,
Salzberg S.L., Fraser C.M., Venter J.C.,
"Sequence and analysis of chromosome 2 of the plant Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae, Streptophyra, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, Rosidae, eurosids II; Brassicales; Brassicaceae, Arabidopsis.
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                                                                                     SEQUENCE FROM N.A.
MEDLINE=95282779; PubMed=7762555;
Sebastio G., Sperandeo M.P., Panico M., de Franchis R., Kraus J.P.
                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                      "The molecular basis of homocystinuria due to cystathionine beta-
synthase deficiency in Italian families, and report of four novel
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Pred. No. 55;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                    'Match 56.0%; Score 28; DB 4; Length 16; Local Similarity 55.6%; Pred. No. 46; es 5; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                             16 AA; 1538 MW; 07455C146B001686 CRC64;
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        Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                           19 AA.
Cystathionine beta-synthase (Fragment) Phomo sapiens (Human)
                                                                                                                                                          mutations.";
Am. J. Hum. Genet. 56:1324-1333(1995)
EMBL; S78267; AAB34404.1;
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Matches 6; Conservative
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STRAIN=CV. COLUMBIA;
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                                                             NCBI_TaxID=9606;
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                       STRAIN-ATCC 15692 / PAO1;
MEDLINE-20437337; PubMed=10984043;
Stover C.K., Pham X.-Q.T., Erwin A.L., Mizoguchi S.D., Warrener P., Stover C.K., Pham X.-Q.T., Erwin A.L., Mizoguchi S.D., Warrener P., Hickey M.J., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagrou M. Garber R.L., Goltry L., Tolentino E., Westbrock-Wadman S., Yuan Y. Brody L.L., Coulter S.N., Folger K.R., Kas A., Larbig K., Lim R.M., Smith K.A., Spencer D.H., Wong G.K.-S., Wu Z., Paulsen I.T., Reizer J., Saier M.H., Hancock R.E.W., Lory S., Olson M.V.; "Complete genome sequence of Pseudomonas aeruginosa PAO1, an opportunistic pathogen.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Bacteria, Proteobacteria, gamma subdivision, Enterobacteriaceae,
                                                                                                                                                PA3577.
Pseudomonas aeruginosa.
Bacteria: Proteobacteria; gamma subdivision; Pseudomonadaceae;
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Martin I.B., Bacon D.J., Tyler S.D., Munro C.K., Johnson W.M.;
J. Clin. Microbiol. 0:0-0(1995).
EMBL; U41250; AAB40560.1; --
HSSP; P08027; IBOV.
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Pred. No. 1.8e+02;
1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nature 406:959-964 (2000).
EMBL, AE004778; AAG06965.1; -.
Hypothetical protein; Complete proteome.
SEQUENCE 37 AA; 4372 MW; 0C68B97A779C241E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Variant shigalike toxin II VT subunit A (Fragment).
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                                                                            Last sequence update)
Last annotation update)
       37 AA
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     PRT;
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PFdm; PF02258; SLT_beta; 1.
NON TER 1 SEQUENCE 42 AA, 4576 MW; BC
                                               ol-mak-2001 (TrEMBLrel. 16, 01-MAR-2001 (TrEMBLrel. 16, 101-OCT-2001 (TrEMBLrel. 18, 14) Pypothetical protein PA3577.
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62.5%;
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Matches 5; Conservative
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GSGFAEVQ 38
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NCBI_TaxID=562;
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P77064;
O1-FEB-1997 (TrEMBLrel. 02, Cr
O1-FEB-1997 (TrEMBLrel. 02, Le
O1-DEC-2001 (TrEMBLrel. 19, Le
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J. Clin. Microbiol. 0:0-0 (1995).
EMBL; U41260; AAB40570.1; -.
HSSP; P08027; 1BOV.
IncorPro; IPR003189; SIT_beta.
Pfam; PF02258; SIT_beta; 1.
NON_TER
                                                           P77498 PRELIMINARY;
P77498;
C1-FEB-1997 (TrEMBLrel. 02, C1
01-FEB-1997 (TrEMBLrel. 17, Le
01-JUN-2001 (TrEMBLrel. 17, Le
Variant SHIGA-like toxin II V
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SEQUENCE
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InterPro; IPR003189; SLT_beta.
Pfam; PF02258; SLT_beta; 1.
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01-FEB-1997 (TrEMBLrel. 02,
01-DEC-2001 (TrEMBLrel. 19,
Variant shiga-like toxin II
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Escherichia coli.
Bacteria; Proteobacteria;
Escherichia.
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62.5%;
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Pred. No.
2; Mismatc
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Best Local S
Matches 5
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Best Local Similarity
Matches 5; Conserv
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P77065;
P77065;
O1-FEB-1997 (TrEMBLrel. 02, Cr
O1-FEB-1997 (TrEMBLrel. 19, Li
O1-DEC-2001 (TrEMBLrel. 19, Li
Variant shiqa-like toxin II V
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EMBL; U41252; AAB40562.1; -.
HSSP; P08027; 1BOV.
InterPro; IPR003189; SIT_beta.
Pfam; PF02258; SIT_beta; 1.
NON_TER
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Submitted (JAN-1997) to the EMBL
EMBL; U41258; AAB40566.1; -.
EMBL; U41256; AAB40566.1; -.
HSSP; P08027; 1BOV.
InterPro; IPR003189; SLT_beta.
Pfan; PF02258; SLT_beta; 1.
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01-FEB-1997
01-JUN-2001
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NON_TER
SEQUENCE FROM N.A.
Martin I.E., Bacon D.J., Tyler
J. Clin. Microbiol. 0:0-0(1995)
EMBL; U41253; AAB40563.1; -
HSSP; P08027; 1BOV.
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01-FEB-1997 (TrEMBLrel. 02,
01-JUN-2001 (TrEMBLrel. 17,
01-JUN-2011 (TrEMBLrel. 17,
01-Junt SHIGA-like toxin II
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Bacteria, Proteobacteria,
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5; Conservative
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                                                                                                 Tyler S.D.,
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the EMBL/GenBank/DDBJ databases.
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he EMBL/GenBank/DDBJ
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VI subunit A (Fragment)
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VT subunit A (Fragment).
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Pred. No. 2.2e-
2; Mismatches
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                                                                                                                                                                                                                                                               subdivision;
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                                                                                                    Munro
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                                                                                                 C.X.,
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Rhabditidae; Peloderinae; Caenorhabditis.
NCBI_TaxID=6239;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Cyclophilin A (Fragment).
PPIA OR CYPA.
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Mus.
NCBI_TaxID=10090;
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                                                                                                           Score 27; DB 2; Length 44;
Pred. No. 2.2e+02;
2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=BRISTOL N2;
Waterston R.;
"Direct Submission.";
Submitted (UUL-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; U58740; AAB00612.1; -..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Martin J.;
"The sequence of C. elegans cosmid R09H3.";
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
InterPro, IPR003189; SLT_beta.
Pfam, PF02258; SLT_beta; 1.
NON TER SEQUENCE 44 AA, 4851 MW, 6579398F654F550C CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Hypothetical 4.6 kDa protein.
R09H3.2.
Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                 39 AA
                                                                                                                                                          2; Mismatches
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MEDLINE=99069613; PubMed=9851916;
                                                                                                              54.0%;
62.5%;
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                                                                                                           Query Match
Best Local Similarity
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MEDLINE=98282291; PubMed=9616227;

MEDLINE=98282291; PubMed=9616227;

MEDLINE=98282291; PubMed=9616227;

Kalama K.E., Trocha A.K., Jones N.G., Johnson P.R.,

Kalama S.A.;

"Lack of strong immune selection pressure by the immunodominant, HLA-

A*0201-restricted cytotoxic T lymphocyte response in chronic human

immunodeficiency virus-1 infection.";

J. Clin. Invest. 101:2559-2566(1998).

EMBL, AF017861; ARC29148.2;

InterPro; IPR000071; Retrovir_p17.

Pfam; PP00540; Gag_D77; I.

AIDS; Core protein; Polyprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                    STRAIN=129/OLA;
MEDLINE=2042570; PubMed=10964515;
COLGAN J., Asmal M., Luban J.;
"Isolation, characterization and targeted disruption of mouse Ppia:
"Isolation, characterization and targeted disruption of mouse Ppia:
cyclophilin A is not essential for mammalian cell viability.";
Genomics 68:167-178(2000).
HSSP; P05092; 2CPL.
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Pred. No. 3.8e+02;
0, Mismatches 4; Indels
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Last sequence update)
Last annotation update)
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01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29 AA
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                                                                                                                                                                                                                     MGD; MGI:97749; Ppia.
InterPro; IPR002130; CSA_PPIase.
Pfam; PF00160; pro isomerae; 1.
PROSITE; PS50072; CSA_PPIASE_2; 1.
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Best Local Similarity 55.6%;
Matches 5; Conservative
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Best Local Similarity 55.00,

Best Local Si Conservative
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SEQUENCE FROM N.A.
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26 GSSFHRIIP 34
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RESULT 15
Q91X54
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Matches 5
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Best Local Similarity
Matches 5; Conserv
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09IX54;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Major core protein (Fragment).
Possum adenovirus.
Viruses; dabNA viruses, no RNA stage; Adenoviridae; Atadenovirus.
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
STRAIN-SPRAGUE-DAWLEY; TISSUE-BRAIN;
MEDLINE-98325069; PubMed-9660825;
Dufour C., Weinberger R.P., Schevzov G., Jeffrey P.L., Gunning P.;
"Splicing of two internal and four carboxyl-terminal alternative exons "Splicing of two internal and four carboxyl-terminal alternative exons in nonmuscle tropomyosin 5 pre-mRNA is independently regulated during
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AF053360; AAC27291.1; -.
InterPro; ITR000533; Tropomyosin.
Pfam; PF00261; Tropomyosin; 1.
NON_TER 1
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J. Biol. Chem. 273:18547-18555(1998).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=121816;
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                                                                                                                       25 GLRFSKROP 33
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Pred. No. 5.6e-
1; Mismatches
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